-34-

SEQUENCE LISTING

Assignment of SEQ ID NOS

	SEQ ID NO 1	acetoacetyl-CoA	thiolase nucle	otide sequence	e		
5	SEQ ID NO 2	HMG-CoA synth	ase nucleotide	sequence			
	SEQ ID NO 3	HMG-CoA reduc	tase nucleotid	e sequence			
	SEQ ID NO 4	mevalonate kinas	se nucletotide s	sequence			
	SEQ ID NO 5	phosphomevalon	ate kinase nuc	leotide sequer	nce		
	SEQ ID NO 6	mevalonate pyro	phosphate deca	arboxylase nu	cleotide seque	ence	
10	SEQ ID NO 7	"single operon" nucleotide sequence					
	SEQ ID NO 8	"MEVT" operon	nucleotide sec	quence			
	SEQ ID NO 9	"MEVB" operon			ncluding the tr	anscription	
And the second s	terminator)	1		`	_		
	SEQ ID NO 10	Isopentenyl pyro	phosphate ison	merase (<i>idi</i>) n	acleotide sequ	ience	
15	SEQ ID NO 11	Farnesyl pyropho					
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E	SEQ ID NO 12	"MBI" operon nu	icleotide seque	encer			
His miles from the first from the fi	SEQ ID NO 13	"MBIS" operon	nucleotide seq	uence			
The second secon							
plant of the party							
20	SEQ ID NO 1	Acetoacetyl-CoA	A thiolase nucl	eotide sequen	ce		
			CA CITIC COCCEA	COMP OFFICER		ምል እ <i>ርርር</i> ምምር እ	
		AATT GTGTCATCGT					
		rcca ccagcgccat					
		ATCG ATTCACAACA					
25		CAAA ATCCGGCGCG					
		ACGG TCAATAAAGT					
		CAGG CAGGTCAGGC					
		FACT TACTCGATGC					
		GTAA TCCTGCGCGA					
30		GAAA ACGTGGCTAA					
		TCAC AGCGTAAAGC					
	601 GTCCCG	GTAA ATGTTGTCAC	TCGAAAGAAA	ACCTTCGTCT	TCAGTCAAGA	CGAATTCCCG	

661	AAAGCGAATT	CAACGGCTGA	AGCGTTAGGT	GCATTGCGCC	CGGCCTTCGA	TAAAGCAGGA
721	ACAGTCACCG	CTGGGAACGC	GTCTGGTATT	AACGACGGTG	CTGCCGCTCT	GGTGATTATG
781	GAAGAATCTG	CGGCGCTGGC	AGCAGGCCTT	ACCCCCTGG	CTCGCATTAA	AAGTTATGCC
841	AGCGGTGGCG	TGCCCCCGC	ATTGATGGGT	ATGGGGCCAG	TACCTGCCAC	GCAAAAAGCG
901	TTACAACTGG	CGGGGCTGCA	ACTGGCGGAT	ATTGATCTCA	TTGAGGCTAA	TGAAGCATTT
961	GCTGCACAGT	TCCTTGCCGT	TGGGAAAAAC	CTGGGCTTTG	ATTCTGAGAA	AGTGAATGTC
1021	AACGGCGGGG	CCATCGCGCT	CGGGCATCCT	ATCGGTGCCA	GTGGTGCTCG	TATTCTGGTC
1081	ACACTATTAC	ATGCCATGCA	GGCACGCGAT	AAAACGCTGG	GGCTGGCAAC	ACTGTGCATT
1141	GGCGGCGGTC	AGGGAATTGC	GATGGTGATT	GAACGGTTGA	ATTAA	

SEQ ID NO 2 HMG-CoA synthase nucleotide sequence

	1	ATGAAACTCT	CAACTAAACT	TTGTTGGTGT	GGTATTAAAG	GAAGACTTAG	GCCGCAAAAG
	61	CAACAACAAT	TACACAATAC	AAACTTGCAA	ATGACTGAAC	TAAAAAAACA	AAAGACCGCT
1	.21	GAACAAAAAA	CCAGACCTCA	AAATGTCGGT	ATTAAAGGTA	TCCAAATTTA	CATCCCAACT
1	181	CAATGTGTCA	ACCAATCTGA	GCTAGAGAAA	TTTGATGGCG	TTTCTCAAGG	TAAATACACA
2	241	ATTGGTCTGG	GCCAAACCAA	CATGTCTTTT	GTCAATGACA	GAGAAGATAT	CTACTCGATG
3	301	TCCCTAACTG	TTTTGTCTAA	GTTGATCAAG	AGTTACAACA	TCGACACCAA	CAAAATTGGT
3	361	AGATTAGAAG	TCGGTACTGA	AACTCTGATT	GACAAGTCCA	AGTCTGTCAA	GTCTGTCTTG
4	121	ATGCAATTGT	TTGGTGAAAA	CACTGACGTC	GAAGGTATTG	ACACGCTTAA	TGCCTGTTAC
4	181	GGTGGTACCA	ACGCGTTGTT	CAACTCTTTG	AACTGGATTG	AATCTAACGC	ATGGGATGGT
į	541	AGAGACGCCA	TTGTAGTTTG	CGGTGATATT	GCCATCTACG	ATAAGGGTGC	CGCAAGACCA
(601	ACCGGTGGTG	CCGGTACTGT	TGCTATGTGG	ATCGGTCCTG	ATGCTCCAAT	TGTATTTGAC
(661	TCTGTAAGAG	CTTCTTACAT	GGAACACGCC	TACGATTTTT	ACAAGCCAGA	TTTCACCAGC
	721	GAATATCCTT	ACGTCGATGG	TCATTTTTCA	TTAACTTGTT	ACGTCAAGGC	TCTTGATCAA
	781	GTTTACAAGA	GTTATTCCAA	GAAGGCTATT	TCTAAAGGGT	TGGTTAGCGA	TCCCGCTGGT
	841	TCGGATGCTT	TGAACGTTTT	GAAATATTTC	GACTACAACG	TTTTCCATGT	TCCAACCTGT
	901	AAATTGGTCA	CAAAATCATA	CGGTAGATTA	CTATATAACG	ATTTCAGAGC	CAATCCTCAA
	961	TTGTTCCCAG	AAGTTGACGC	CGAATTAGCT	ACTCGCGATT	ATGACGAATC	TTTAACCGAT
1	021	AAGAACATTG	AAAAAACTTT	TGTTAATGTT	GCTAAGCCAT	TCCACAAAGA	GAGAGTTGCC
1	081	CAATCTTTGA	TTGTTCCAAC	AAACACAGGT	AACATGTACA	CCGCATCTGT	TTATGCCGCC
1	141	TTTGCATCTC	TATTAAACTA	TGTTGGATCT	GACGACTTAC	AAGGCAAGCG	TGTTGGTTTA
1	201	TTTTCTTACG	GTTCCGGTTT	AGCTGCATCT	CTATATTCTT	GCAAAATTGT	TGGTGACGTC
1	261	CAACATATTA	TCAAGGAATT	AGATATTACT	AACAAATTAG	CCAAGAGAAT	CACCGAAACT
1	321	CCAAAGGATT	ACGAAGCTGC	CATCGAATTG	AGAGAAAATG	CCCATTTGAA	GAAGAACTTC
1	381	AAACCTCAAG	GTTCCATTGA	GCATTTGCAA	AGTGGTGTT	ACTACTTGAC	CAACATCGAT

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1441 GACAAATTTA GAAGATCTTA CGATGTTAAA AAATAA

SEQ ID NO 3 Truncated HMG-CoA reductase nucleotide sequence

Artificial Start Codon

1 ATGGTTTTAA CCAATAAAAC AGTCATTTCT GGATCGAAAG TCAAAAGTTT ATCATCTGCG
Base Pair 1660 of S.cervisiae HMG1

61 CAATCGAGCT CATCAGGACC TTCATCATCT AGTGAGGAAG ATGATTCCCG CGATATTGAA 121 AGCTTGGATA AGAAAATACG TCCTTTAGAA GAATTAGAAG CATTATTAAG TAGTGGAAAT 181 ACAAAACAAT TGAAGAACAA AGAGGTCGCT GCCTTGGTTA TTCACGGTAA GTTACCTTTG 241 TACGCTTTGG AGAAAAAATT AGGTGATACT ACGAGAGCGG TTGCGGTACG TAGGAAGGCT 301 CTTTCAATTT TGGCAGAAGC TCCTGTATTA GCATCTGATC GTTTACCATA TAAAAATTAT 361 GACTACGACC GCGTATTTGG CGCTTGTTGT GAAAATGTTA TAGGTTACAT GCCTTTGCCC 421 GTTGGTGTTA TAGGCCCCTT GGTTATCGAT GGTACATCTT ATCATATACC AATGGCAACT 481 ACAGAGGGTT GTTTGGTAGC TTCTGCCATG CGTGGCTGTA AGGCAATCAA TGCTGGCGGT 541 GGTGCAACAA CTGTTTTAAC TAAGGATGGT ATGACAAGAG GCCCAGTAGT CCGTTTCCCA 601 ACTTTGAAAA GATCTGGTGC CTGTAAGATA TGGTTAGACT CAGAAGAGGG ACAAAACGCA 661 ATTAAAAAG CTTTTAACTC TACATCAAGA TTTGCACGTC TGCAACATAT TCAAACTTGT 721 CTAGCAGGAG ATTTACTCTT CATGAGATTT AGAACAACTA CTGGTGACGC AATGGGTATG 781 AATATGATTT CTAAAGGTGT CGAATACTCA TTAAAGCAAA TGGTAGAAGA GTATGGCTGG 841 GAAGATATGG AGGTTGTCTC CGTTTCTGGT AACTACTGTA CCGACAAAAA ACCAGCTGCC 901 ATCAACTGGA TCGAAGGTCG TGGTAAGAGT GTCGTCGCAG AAGCTACTAT TCCTGGTGAT 961 GTTGTCAGAA AAGTGTTAAA AAGTGATGTT TCCGCATTGG TTGAGTTGAA CATTGCTAAG 1021 AATTTGGTTG GATCTGCAAT GGCTGGGTCT GTTGGTGGAT TTAACGCACA TGCAGCTAAT 1081 TTAGTGACAG CTGTTTTCTT GGCATTAGGA CAAGATCCTG CACAAAATGT TGAAAGTTCC 1141 AACTGTATAA CATTGATGAA AGAAGTGGAC GGTGATTTGA GAATTTCCGT ATCCATGCCA 1201 TCCATCGAAG TAGGTACCAT CGGTGGTGGT ACTGTTCTAG AACCACAAGG TGCCATGTTG 1261 GACTTATTAG GTGTAAGAGG CCCGCATGCT ACCGCTCCTG GTACCAACGC ACGTCAATTA 1321 GCAAGAATAG TTGCCTGTGC CGTCTTGGCA GGTGAATTAT CCTTATGTGC TGCCCTAGCA 1381 GCCGGCCATT TGGTTCAAAG TCATATGACC CACAACAGGA AACCTGCTGA ACCAACAAAA 1441 CCTAACAATT TGGACGCCAC TGATATAAAT CGTTTGAAAG ATGGGTCCGT CACCTGCATT 1501 AAATCCTAA

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SEQ ID NO 4 Mevalonate kinase nucletotide sequence

1	ATGTCATTAC	CGTTCTTAAC	TTCTGCACCG	GGAAAGGTTA	TTATTTTTGG	TGAACACTCT
61	GCTGTGTACA	ACAAGCCTGC	CGTCGCTGCT	AGTGTGTCTG	CGTTGAGAAC	CTACCTGCTA
121	ATAAGCGAGT	CATCTGCACC	AGATACTATT	GAATTGGACT	TCCCGGACAT	TAGCTTTAAT
181	CATAAGTGGT	CCATCAATGA	TTTCAATGCC	ATCACCGAGG	ATCAAGTAAA	CTCCCAAAAA
241	TTGGCCAAGG	CTCAACAAGC	CACCGATGGC	TTGTCTCAGG	AACTCGTTAG	TCTTTTGGAT
301	CCGTTGTTAG	CTCAACTATC	CGAATCCTTC	CACTACCATG	CAGCGTTTTG	TTTCCTGTAT
361	ATGTTTGTTT	GCCTATGCCC	CCATGCCAAG	AATATTAAGT	TTTCTTTAAA	GTCTACTTTA
421	CCCATCGGTG	CTGGGTTGGG	CTCAAGCGCC	TCTATTTCTG	TATCACTGGC	CTTAGCTATG
481	GCCTACTTGG	GGGGGTTAAT	AGGATCTAAT	GACTTGGAAA	AGCTGTCAGA	AAACGATAAG
541	CATATAGTGA	ATCAATGGGC	CTTCATAGGT	GAAAAGTGTA	TTCACGGTAC	CCCTTCAGGA
601	ATAGATAACG	CTGTGGCCAC	TTATGGTAAT	GCCCTGCTAT	TTGAAAAAGA	CTCACATAAT
661	GGAACAATAA	ACACAAACAA	TTTTAAGTTC	TTAGATGATT	TCCCAGCCAT	TCCAATGATC
721	CTAACCTATA	CTAGAATTCC	AAGGTCTACA	AAAGATCTTG	TTGCTCGCGT	TCGTGTGTTG
781	GTCACCGAGA	AATTTCCTGA	AGTTATGAAG	CCAATTCTAG	ATGCCATGGG	TGAATGTGCC
841	CTACAAGGCT	TAGAGATCAT	GACTAAGTTA	AGTAAATGTA	AAGGCACCGA	TGACGAGGCT
901	GTAGAAACTA	ATAATGAACT	GTATGAACAA	CTATTGGAAT	TGATAAGAAT	AAATCATGGA
961	CTGCTTGTCT	CAATCGGTGT	TTCTCATCCT	GGATTAGAAC	TTATTAAAAA	TCTGAGCGAT
1021	GATTTGAGAA	TTGGCTCCAC	AAAACTTACC	GGTGCTGGTG	GCGGCGGTTG	CTCTTTGACT
1081	TTGTTACGAA	GAGACATTAC	TCAAGAGCAA	ATTGACAGCT	TCAAAAAGAA	ATTGCAAGAT
1141	GATTTTAGTT	ACGAGACATT	TGAAACAGAC	TTGGGTGGGA	CTGGCTGCTG	TTTGTTAAGC
1201	GCAAAAAATT	TGAATAAAGA	TCTTAAAATC	AAATCCCTAG	TATTCCAATT	ATTTGAAAAT
1261	AAAACTACCA	CAAAGCAACA	AATTGACGAT	CTATTATTGC	CAGGAAACAC	GAATTTACCA
1321	TGGACTTCAT	AG				

SEQ ID NO 5 Phosphomevalonate kinase nucleotide sequence

1	ATGTCAGAGT	TGAGAGCCTT	CAGTGCCCCA	GGGAAAGCGT	TACTAGCTGG	TGGATATTTA
61	GTTTTAGATA	CAAAATATGA	AGCATTTGTA	GTCGGATTAT	CGGCAAGAAT	GCATGCTGTA
121	GCCCATCCTT	ACGGTTCATT	GCAAGGGTCT	GATAAGTTTG	AAGTGCGTGT	GAAAAGTAAA
181	CAATTTAAAG	ATGGGGAGTG	GCTGTACCAT	ATAAGTCCTA	AAAGTGGCTT	CATTCCTGTT
241	TCGATAGGCG	GATCTAAGAA	CCCTTTCATT	GAAAAGTTA	TCGCTAACGT	ATTTAGCTAC
301	TTTAAACCTA	ACATGGACGA	CTACTGCAAT	AGAAACTTGT	TCGTTATTGA	TATTTTCTCT
361	GATGATGCCT	ACCATTCTCA	GGAGGATAGC	GTTACCGAAC	ATCGTGGCAA	CAGAAGATTG
421	AGTTTTCATT	CGCACAGAAT	TGAAGAAGTT	CCCAAAACAG	GGCTGGGCTC	CTCGGCAGGT
481	TTAGTCACAG	TTTTAACTAC	AGCTTTGGCC	TCCTTTTTTG	TATCGGACCT	GGAAAATAAT
541	GTAGACAAAT	ATAGAGAAGT	TATTCATAAT	TTAGCACAAG	TTGCTCATTG	TCAAGCTCAG

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601	GGTAAAATTG	GAAGCGGGTT	TGATGTAGCG	GCGGCAGCAT	ATGGATCTAT	CAGATATAGA
661	AGATTCCCAC	CCGCATTAAT	CTCTAATTTG	CCAGATATTG	GAAGTGCTAC	TTACGGCAGT
721	AAACTGGCGC	ATTTGGTTGA	TGAAGAAGAC	TGGAATATTA	CGATTAAAAG	TAACCATTTA
781	CCTTCGGGAT	TAACTTTATG	GATGGGCGAT	ATTAAGAATG	GTTCAGAAAC	AGTAAAACTG
841	GTCCAGAAGG	TAAAAAATTG	GTATGATTCG	CATATGCCAG	AAAGCTTGAA	AATATATACA
901	GAACTCGATC	ATGCAAATTC	TAGATTTATG	GATGGACTAT	CTAAACTAGA	TCGCTTACAC
961	GAGACTCATG	ACGATTACAG	CGATCAGATA	TTTGAGTCTC	TTGAGAGGAA	TGACTGTACC
1021	TGTCAAAAGT	ATCCTGAAAT	CACAGAAGTT	AGAGATGCAG	TTGCCACAAT	TAGACGTTCC
1081	TTTAGAAAAA	TAACTAAAGA	ATCTGGTGCC	GATATCGAAC	CTCCCGTACA	AACTAGCTTA
1141	TTGGATGATT	GCCAGACCTT	AAAAGGAGTT	CTTACTTGCT	TAATACCTGG	TGCTGGTGGT
1201	TATGACGCCA	TTGCAGTGAT	TACTAAGCAA	GATGTTGATC	TTAGGGCTCA	AACCGCTAAT
1261	GACAAAAGAT	TTTCTAAGGT	TCAATGGCTG	GATGTAACTC	AGGCTGACTG	GGGTGTTAGG
1321	AAAGAAAAAG	ATCCGGAAAC	TTATCTTGAT	AAATAG		
SEQ ID NO) 6 Me	evalonate pyrc	phosphate de	carboxylase n	ucleotide sequ	ience
1	ATGACCGTTT	ACACAGCATC	CGTTACCGCA	CCCGTCAACA	TCGCAACCCT	TAAGTATTGG
61	GGGAAAAGGG	ACACGAAGTT	GAATCTGCCC	ACCAATTCGT	CCATATCAGT	GACTTTATCG
121	CAAGATGACC	TCAGAACGTT	GACCTCTGCG	GCTACTGCAC	CTGAGTTTGA	ACGCGACACT
181	TTGTGGTTAA	ATGGAGAACC	ACACAGCATC	GACAATGAAA	GAACTCAAAA	TTGTCTGCGC
241	GACCTACGCC	AATTAAGAAA	GGAAATGGAA	TCGAAGGACG	CCTCATTGCC	CACATTATCT
301	CAATGGAAAC	TCCACATTGT	CTCCGAAAAT	AACTTTCCTA	CAGCAGCTGG	TTTAGCTTCC
361	TCCGCTGCTG	GCTTTGCTGC	ATTGGTCTCT	GCAATTGCTA	AGTTATACCA	ATTACCACAG
421	TCAACTTCAG	AAATATCTAG	AATAGCAAGA	AAGGGGTCTG	GTTCAGCTTG	TAGATCGTTG
481	TTTGGCGGAT	ACGTGGCCTG	GGAAATGGGA	AAAGCTGAAG	ATGGTCATGA	TTCCATGGCA
541	GTACAAATCG	CAGACAGCTC	TGACTGGCCT	CAGATGAAAG	CTTGTGTCCT	AGTTGTCAGC
601	GATATTAAAA	AGGATGTGAG	TTCCACTCAG	GGTATGCAAT	TGACCGTGGC	AACCTCCGAA
661	CTATTTAAAG	AAAGAATTGA	ACATGTCGTA	CCAAAGAGAT	TTGAAGTCAT	GCGTAAAGCC
721	ATTGTTGAAA	AAGATTTCGC	CACCTTTGCA	AAGGAAACAA	TGATGGATTC	CAACTCTTTC
781	CATGCCACAT	GTTTGGACTC	TTTCCCTCCA	ATATTCTACA	TGAATGACAC	TTCCAAGCGT
841	ATCATCAGTT	' GGTGCCACAC	CATTAATCAG	TTTTACGGAG	AAACAATCGT	TGCATACACG
901	TTTGATGCAG	GTCCAAATGC	TGTGTTGTAC	TACTTAGCTG	AAAATGAGTC	GAAACTCTTT
961	GCATTTATCT	' ATAAATTGTT	TGGCTCTGTT	CCTGGATGGG	ACAAGAAATT	TACTACTGAG
						TGAATTGGAT
1081	CTTGAGTTGC	AAAAGGATGT	TGCCAGAGTG	ATTTTAACTC	: AAGTCGGTTC	: AGGCCCACAA

1141 GAAACAAACG AATCTTTGAT TGACGCAAAG ACTGGTCTAC CAAAGGAATA A

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SEQ ID NO 7 "single operon" nucleotide sequence

P_{BAD} Promoter

1 GACGCTTTTT ATCGCAACTC TCTACTGTTT CTCCATACCC GTTTTTTTGG GCTAGCAGGA

RBS Start Isopentyl
61 GGAATTCACC ATGGTACCCG GGAGGAGT TACTATATGC AAACGGAACA CGTCATTTTA

Pyrophosphate Isomerase

TTGAATGCAC AGGGAGTTCC CACGGGTACG CTGGAAAAGT ATGCCGCACA CACGGCAGAC

181 ACCCGCTTAC ATCTCGCGTT CTCCAGTTGG CTGTTTAATG CCAAAGGACA ATTATTAGTT

241 ACCCGCCGCG CACTGAGCAA AAAAGCATGG CCTGGCGTGT GGACTAACTC GGTTTGTGGG

301 CACCCACAAC TGGGAGAAAG CAACGAAGAC GCAGTGATCC GCCGTTGCCG TTATGAGCTT

361 GGCGTGGAAA TTACGCCTCC TGAATCTATC TATCCTGACT TTCGCTACCG CGCCACCGAT

421 CCGAGTGGCA TTGTGGAAAA TGAAGTGTGT CCGGTATTTG CCGCACGCAC CACTAGTGCG

481 TTACAGATCA ATGATGATGA AGTGATGGAT TATCAATGGT GTGATTAGC AGATGTATTA

541 CACGGTATTG ATGCCACGCC GTGGGCGTTC AGTCCGTGGA TGGTGATGCA GGCGACAAAT

End Isopentyl Pyrophosphate Isomerase

601 CGCGAAGCCA GAAAACGATT ATCTGCATTT ACCCAGCTTA AATAACCCGG GGATCCTCTA

Start Acetoacetyl-CoA Thiolase RBS 661 GAGTCGACTA GGAGGAATAT AAAATGAAAA ATTGTGTCAT CGTCAGTGCG GTACGTACTG 721 CTATCGGTAG TTTTAACGGT TCACTCGCTT CCACCAGCGC CATCGACCTG GGGGCGACAG 781 TAATTAAAGC CGCCATTGAA CGTGCAAAAA TCGATTCACA ACACGTTGAT GAAGTGATTA 841 TGGGTAACGT GTTACAAGCC GGGCTGGGGC AAAATCCGGC GCGTCAGGCA CTGTTAAAAA 901 GCGGGCTGGC AGAAACGGTG TGCGGATTCA CGGTCAATAA AGTATGTGGT TCGGGTCTTA 961 AAAGTGTGGC GCTTGCCGCC CAGGCCATTC AGGCAGGTCA GGCGCAGAGC ATTGTGGCGG 1021 GGGGTATGGA AAATATGAGT TTAGCCCCCT ACTTACTCGA TGCAAAAGCA CGCTCTGGTT 1081 ATCGTCTTGG AGACGGACAG GTTTATGACG TAATCCTGCG CGATGGCCTG ATGTGCGCCA 1141 CCCATGGTTA TCATATGGGG ATTACCGCCG AAAACGTGGC TAAAGAGTAC GGAATTACCC 1201 GTGAAATGCA GGATGAACTG GCGCTACATT CACAGCGTAA AGCGGCAGCC GCAATTGAGT 1261 CCGGTGCTTT TACAGCCGAA ATCGTCCCGG TAAATGTTGT CACTCGAAAG AAAACCTTCG 1321 TCTTCAGTCA AGACGAATTC CCGAAAGCGA ATTCAACGGC TGAAGCGTTA GGTGCATTGC 1381 GCCCGGCCTT CGATAAAGCA GGAACAGTCA CCGCTGGGAA CGCGTCTGGT ATTAACGACG 1441 GTGCTGCCGC TCTGGTGATT ATGGAAGAAT CTGCGGCGCT GGCAGCAGGC CTTACCCCCC

1501 TGGCTCGCAT TAAAAGTTAT GCCAGCGGTG GCGTGCCCCC CGCATTGATG GGTATGGGGC 1561 CAGTACCTGC CACGCAAAAA GCGTTACAAC TGGCGGGGCT GCAACTGGCG GATATTGATC 1621 TCATTGAGGC TAATGAAGCA TTTGCTGCAC AGTTCCTTGC CGTTGGGAAA AACCTGGGCT 1681 TTGATTCTGA GAAAGTGAAT GTCAACGGCG GGGCCATCGC GCTCGGGCAT CCTATCGGTG 1741 CCAGTGGTGC TCGTATTCTG GTCACACTAT TACATGCCAT GCAGGCACGC GATAAAACGC 1801 TGGGGCTGGC AACACTGTGC ATTGGCGGCG GTCAGGGAAT TGCGATGGTG ATTGAACGGT Stop Acetoacetyl-CoA Start HMG-CoA Synthase Thiolase____ 1861 TGAATTAAGG AGGACAGCTA AATGAAACTC TCAACTAAAC TTTGTTGGTG TGGTATTAAA 1921 GGAAGACTTA GGCCGCAAAA GCAACAACAA TTACACAATA CAAACTTGCA AATGACTGAA 1981 CTAAAAAAC AAAAGACCGC TGAACAAAAA ACCAGACCTC AAAATGTCGG TATTAAAGGT 2041 ATCCAAATTT ACATCCCAAC TCAATGTGTC AACCAATCTG AGCTAGAGAA ATTTGATGGC 2101 GTTTCTCAAG GTAAATACAC AATTGGTCTG GGCCAAACCA ACATGTCTTT TGTCAATGAC 2161 AGAGAAGATA TCTACTCGAT GTCCCTAACT GTTTTGTCTA AGTTGATCAA GAGTTACAAC 2221 ATCGACACCA ACAAAATTGG TAGATTAGAA GTCGGTACTG AAACTCTGAT TGACAAGTCC 2281 AAGTCTGTCA AGTCTGTCTT GATGCAATTG TTTGGTGAAA ACACTGACGT CGAAGGTATT 2341 GACACGCTTA ATGCCTGTTA CGGTGGTACC AACGCGTTGT TCAACTCTTT GAACTGGATT 2401 GAATCTAACG CATGGGATGG TAGAGACGCC ATTGTAGTTT GCGGTGATAT TGCCATCTAC 2461 GATAAGGGTG CCGCAAGACC AACCGGTGGT GCCGGTACTG TTGCTATGTG GATCGGTCCT 2521 GATGCTCCAA TTGTATTTGA CTCTGTAAGA GCTTCTTACA TGGAACACGC CTACGATTTT 2581 TACAAGCCAG ATTTCACCAG CGAATATCCT TACGTCGATG GTCATTTTTC ATTAACTTGT 2641 TACGTCAAGG CTCTTGATCA AGTTTACAAG AGTTATTCCA AGAAGGCTAT TTCTAAAGGG 2701 TTGGTTAGCG ATCCCGCTGG TTCGGATGCT TTGAACGTTT TGAAATATTT CGACTACAAC 2761 GTTTTCCATG TTCCAACCTG TAAATTGGTC ACAAAATCAT ACGGTAGATT ACTATATAAC 2821 GATTTCAGAG CCAATCCTCA ATTGTTCCCA GAAGTTGACG CCGAATTAGC TACTCGCGAT 2881 TATGACGAAT CTTTAACCGA TAAGAACATT GAAAAAACTT TTGTTAATGT TGCTAAGCCA 2941 TTCCACAAAG AGAGAGTTGC CCAATCTTTG ATTGTTCCAA CAAACACAGG TAACATGTAC 3001 ACCGCATCTG TTTATGCCGC CTTTGCATCT CTATTAAACT ATGTTGGATC TGACGACTTA 3061 CAAGGCAAGC GTGTTGGTTT ATTTTCTTAC GGTTCCGGTT TAGCTGCATC TCTATATTCT 3121 TGCAAAATTG TTGGTGACGT CCAACATATT ATCAAGGAAT TAGATATTAC TAACAAATTA 3181 GCCAAGAGAA TCACCGAAAC TCCAAAGGAT TACGAAGCTG CCATCGAATT GAGAGAAAAT 3241 GCCCATTTGA AGAAGAACTT CAAACCTCAA GGTTCCATTG AGCATTTGCA AAGTGGTGTT

Stop HMG-CoA synthase____

3301 TACTACTTGA CCAACATCGA TGACAAATTT AGAAGATCTT ACGATGTTAA AAAATAAGGA

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RBS

Start Truncated HMG-CoA Reductase

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	3361 GGATTACACT ATGGTTTTAA CCAATAAAAC AGTCATTTCT GGATCGAAAG TCAAAAGTTT
5	3421 ATCATCTGCG CAATCGAGCT CATCAGGACC TTCATCATCT AGTGAGGAAG ATGATTCCCG
	3481 CGATATTGAA AGCTTGGATA AGAAAATACG TCCTTTAGAA GAATTAGAAG CATTATTAAG
	3541 TAGTGGAAAT ACAAAACAAT TGAAGAACAA AGAGGTCGCT GCCTTGGTTA TTCACGGTAA
	3601 GTTACCTTTG TACGCTTTGG AGAAAAAATT AGGTGATACT ACGAGAGCGG TTGCGGTACG
	3661 TAGGAAGGCT CTTTCAATTT TGGCAGAAGC TCCTGTATTA GCATCTGATC GTTTACCATA
10	3721 TAAAAATTAT GACTACGACC GCGTATTTGG CGCTTGTTGT GAAAATGTTA TAGGTTACAT
	3781 GCCTTTGCCC GTTGGTGTTA TAGGCCCCTT GGTTATCGAT GGTACATCTT ATCATATACC
2 3.	3841 AATGGCAACT ACAGAGGGTT GTTTGGTAGC TTCTGCCATG CGTGGCTGTA AGGCAATCAA
Company of the Compan	3901 TGCTGGCGGT GGTGCAACAA CTGTTTTAAC TAAGGATGGT ATGACAAGAG GCCCAGTAGT
productives and the contract productive in the contract	3961 CCGTTTCCCA ACTTTGAAAA GATCTGGTGC CTGTAAGATA TGGTTAGACT CAGAAGAGGG
15 T	4021 ACAAAACGCA ATTAAAAAAG CTTTTAACTC TACATCAAGA TTTGCACGTC TGCAACATAT
To the second se	4081 TCAAACTTGT CTAGCAGGAG ATTTACTCTT CATGAGATTT AGAACAACTA CTGGTGACGC
The state of the s	4141 AATGGGTATG AATATGATTT CTAAAGGTGT CGAATACTCA TTAAAGCAAA TGGTAGAAGA
A STATE OF S	4201 GTATGGCTGG GAAGATATGG AGGTTGTCTC CGTTTCTGGT AACTACTGTA CCGACAAAAA
12 II	4261 ACCAGCTGCC ATCAACTGGA TCGAAGGTCG TGGTAAGAGT GTCGTCGCAG AAGCTACTAT
20	4321 TCCTGGTGAT GTTGTCAGAA AAGTGTTAAA AAGTGATGTT TCCGCATTGG TTGAGTTGAA
TOTAL	4381 CATTGCTAAG AATTTGGTTG GATCTGCAAT GGCTGGGTCT GTTGGTGGAT TTAACGCACA
International Conference of the Conference of th	4441 TGCAGCTAAT TTAGTGACAG CTGTTTTCTT GGCATTAGGA CAAGATCCTG CACAAAATGT
	4501 TGAAAGTTCC AACTGTATAA CATTGATGAA AGAAGTGGAC GGTGATTTGA GAATTTCCGT
	4561 ATCCATGCCA TCCATCGAAG TAGGTACCAT CGGTGGTGGT ACTGTTCTAG AACCACAAGG
25	4621 TGCCATGTTG GACTTATTAG GTGTAAGAGG CCCGCATGCT ACCGCTCCTG GTACCAACGC
	4681 ACGTCAATTA GCAAGAATAG TTGCCTGTGC CGTCTTGGCA GGTGAATTAT CCTTATGTGC
	4741 TGCCCTAGCA GCCGGCCATT TGGTTCAAAG TCATATGACC CACAACAGGA AACCTGCTGA
	4801 ACCAACAAAA CCTAACAATT TGGACGCCAC TGATATAAAT CGTTTGAAAG ATGGGTCCGT
	Stop Truncated
30	HMG-CoA Reductase RBS Start
	4861 CACCTGCATT AAATCCTAAG TCGACCTGCA GTAGGAGGAA TTAACCATGT CATTACCGTT
	Mevalonate Kinase
	4921 CTTAACTTCT GCACCGGGAA AGGTTATTAT TTTTGGTGAA CACTCTGCTG TGTACAACAA
35	4981 GCCTGCCGTC GCTGCTAGTG TGTCTGCGTT GAGAACCTAC CTGCTAATAA GCGAGTCATC

5041 TGCACCAGAT ACTATTGAAT TGGACTTCCC GGACATTAGC TTTAATCATA AGTGGTCCAT

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51	101	CAATGATTTC	AATGCCATCA	CCGAGGATCA	AGTAAACTCC	CAAAAATTGG	CCAAGGCTCA
51	161	ACAAGCCACC	GATGGCTTGT	CTCAGGAACT	CGTTAGTCTT	TTGGATCCGT	TGTTAGCTCA
52	221	ACTATCCGAA	TCCTTCCACT	ACCATGCAGC	GTTTTGTTTC	CTGTATATGT	TTGTTTGCCT
52	281	ATGCCCCCAT	GCCAAGAATA	TTAAGTTTTC	TTTAAAGTCT	ACTTTACCCA	TCGGTGCTGG
53	341	GTTGGGCTCA	AGCGCCTCTA	TTTCTGTATC	ACTGGCCTTA	GCTATGGCCT	ACTTGGGGGG
54	401	GTTAATAGGA	TCTAATGACT	TGGAAAAGCT	GTCAGAAAAC	GATAAGCATA	TAGTGAATCA
54	461	ATGGGCCTTC	ATAGGTGAAA	AGTGTATTCA	CGGTACCCCT	TCAGGAATAG	ATAACGCTGT
55	521	GGCCACTTAT	GGTAATGCCC	TGCTATTTGA	AAAAGACTCA	CATAATGGAA	CAATAAACAC
55	581	AAACAATTTT	AAGTTCTTAG	ATGATTTCCC	AGCCATTCCA	ATGATCCTAA	CCTATACTAG
56	641	AATTCCAAGG	TCTACAAAAG	ATCTTGTTGC	TCGCGTTCGT	GTGTTGGTCA	CCGAGAAATT
5	701	TCCTGAAGTT	ATGAAGCCAA	TTCTAGATGC	CATGGGTGAA	TGTGCCCTAC	AAGGCTTAGA
5	761	GATCATGACT	AAGTTAAGTA	AATGTAAAGG	CACCGATGAC	GAGGCTGTAG	AAACTAATAA
5	821	TGAACTGTAT	GAACAACTAT	TGGAATTGAT	AAGAATAAAT	CATGGACTGC	TTGTCTCAAT
5	881	CGGTGTTTCT	CATCCTGGAT	TAGAACTTAT	TAAAAATCTG	AGCGATGATT	TGAGAATTGG
5	941	CTCCACAAAA	CTTACCGGTG	CTGGTGGCGG	CGGTTGCTCT	TTGACTTTGT	TACGAAGAGA
6	001	CATTACTCAA	GAGCAAATTG	ACAGCTTCAA	AAAGAAATTG	CAAGATGATT	TTAGTTACGA
6	061	GACATTTGAA	ACAGACTTGG	GTGGGACTGG	CTGCTGTTTG	TTAAGCGCAA	AAAATTTGAA
6	121	TAAAGATCTT	AAAATCAAAT	CCCTAGTATT	CCAATTATTT	GAAAATAAAA	CTACCACAAA
					Stop Me	valonate Ki	nase
6	181	GCAACAAATT	GACGATCTAT	TATTGCCAGG	AAACACGAAT	TTACCATGGA	CTTCATAGGA
							RBS
			├── > Start	Phosphomev	alonate Kin	ase	RBS
6	5241	<u>@</u> GCAGATCAA	1 -		alonate Kin		
		100.13	ATGTCAGAGT	TGAGAGCCTT		GGGAAAGCGT	TACTAGCTGG
6	301	TGGATATTTA	ATGTCAGAGT	TGAGAGCCTT CAAAATATGA	CAGTGCCCCA	GGGAAAGCGT GTCGGATTAT	TACTAGCTGG CGGCAAGAAT
6 6	301 361	TGGATATTTA GCATGCTGTA	ATGTCAGAGT GTTTTAGATA GCCCATCCTT	TGAGAGCCTT CAAAATATGA ACGGTTCATT	CAGTGCCCCA AGCATTTGTA	GGGAAAGCGT GTCGGATTAT GATAAGTTTG	TACTAGCTGG CGGCAAGAAT AAGTGCGTGT
6 6 6	5301 5361 5421 5481	TGGATATTTA GCATGCTGTA GAAAAGTAAA CATTCCTGTT	ATGTCAGAGT GTTTTAGATA GCCCATCCTT CAATTTAAAG TCGATAGGCG	TGAGAGCCTT CAAAATATGA ACGGTTCATT ATGGGGAGTG GATCTAAGAA	CAGTGCCCA AGCATTTGTA GCAAGGGTCT GCTGTACCAT CCCTTTCATT	GGGAAAGCGT GTCGGATTAT GATAAGTTTG ATAAGTCCTA GAAAAAGTTA	TACTAGCTGG CGGCAAGAAT AAGTGCGTGT AAAGTGGCTT TCGCTAACGT
6 6 6	5301 5361 5421 5481	TGGATATTTA GCATGCTGTA GAAAAGTAAA CATTCCTGTT	ATGTCAGAGT GTTTTAGATA GCCCATCCTT CAATTTAAAG TCGATAGGCG	TGAGAGCCTT CAAAATATGA ACGGTTCATT ATGGGGAGTG GATCTAAGAA	CAGTGCCCA AGCATTTGTA GCAAGGGTCT GCTGTACCAT CCCTTTCATT	GGGAAAGCGT GTCGGATTAT GATAAGTTTG ATAAGTCCTA GAAAAAGTTA	TACTAGCTGG CGGCAAGAAT AAGTGCGTGT AAAGTGGCTT
6 6 6 6	5301 5361 5421 5481 5541	TGGATATTTA GCATGCTGTA GAAAAGTAAA CATTCCTGTT ATTTAGCTAC TATTTTCTCT	ATGTCAGAGT GTTTTAGATA GCCCATCCTT CAATTTAAAG TCGATAGGCG TTTAAACCTA	TGAGAGCCTT CAAAATATGA ACGGTTCATT ATGGGGAGTG GATCTAAGAA ACATGGACGA ACCATTCTCA	CAGTGCCCA AGCATTTGTA GCAAGGGTCT GCTGTACCAT CCCTTTCATT CTACTGCAAT GGAGGATAGC	GGGAAAGCGT GTCGGATTAT GATAAGTTCCTA GAAAAAGTTA AGAAACTTGT	TACTAGCTGG CGGCAAGAAT AAGTGCGTGT AAAGTGGCTT TCGCTAACGT TCGTTATTGA ATCGTGGCAA
6 6 6 6	5301 5361 5421 5481 5541 5601	TGGATATTTA GCATGCTGTA GAAAAGTAAA CATTCCTGTT ATTTAGCTAC TATTTTCTCT CAGAAGATTC	ATGTCAGAGT GTTTTAGATA GCCCATCCTT CAATTTAAAG TCGATAGGCG TTTAAACCTA GATGATGCCT GATGATGCCT	TGAGAGCCTT CAAAATATGA ACGGTTCATT ATGGGGAGTG GATCTAAGAA ACATGGACGA ACATGCACGACGACCCCCCCCCC	CAGTGCCCA AGCATTTGTA GCAAGGGTCT GCTGTACCAT CCCTTTCATT CTACTGCAAT GGAGGATAGC TGAAGAAGTT	GGGAAAGCGT GTCGGATTAT GATAAGTTCTA GAAAAAGTTA AGAAACTTGT GTTACCGAAC	TACTAGCTGG CGGCAAGAAT AAGTGCGTGT AAAGTGGCTT TCGCTAACGT TCGTTATTGA ATCGTGGCAA GGCTGGGCTC
6 6 6 6 6	5301 5361 5421 5481 5541 5601 5661	TGGATATTTA GCATGCTGTA GAAAAGTAAA CATTCCTGTT ATTTAGCTAC TATTTTCTCT CAGAAGATTC	ATGTCAGAGT GTTTTAGATA GCCCATCCTT CAATTTAAAG TCGATAGGCG TTTAAACCTA GATGATGCCT AGTTTTCATT	TGAGAGCCTT CAAAATATGA ACGGTTCATT ATGGGGAGTG GATCTAAGAA ACATGGACGA CACCATTCTCA CCGCACAGAAT TTTTAACTAC	CAGTGCCCA AGCATTTGTA GCAAGGGTCT GCTGTACCAT CCCTTTCATT CTACTGCAAT GGAGGATAGC TGAAGAAGTT AGCTTTGGCC	GGGAAAGCGT GTCGGATTAT GATAAGTCCTA GAAAAAGTTAT AGAAAACTTGT CCCCAAAACAG	TACTAGCTGG CGGCAAGAAT AAGTGCGTGT AAAGTGGCTT TCGCTAACGT TCGTTATTGA ATCGTGGCAA GGCTGGGCTC TATCGGACCT
6 6 6 6 6 6	5301 5361 5421 5481 5541 6601 6661 6721	TGGATATTTA GCATGCTGTA GAAAAGTAAA CATTCCTGTT ATTTAGCTAC TATTTTCTCT CAGAAGATTC CTCGGCAGGT	ATGTCAGAGT GTTTTAGATA GCCCATCCTT CAATTTAAAG TCGATAGGCG TTTAAACCTA GATGATGCCT GATGATGCCT TTAGTCACTA TTAGTCACAG	TGAGAGCCTT CAAAATATGA ACGGTTCATT ATGGGGAGTG GATCTAAGAA ACATGGACGA ACCATTCTCA CGCACAGAAT TTTTAACTAC ATAGAGAAGT	CAGTGCCCA AGCATTTGTA GCAAGGGTCT GCTGTACCAT CCCTTTCATT CTACTGCAAT GGAGGATAGC TGAAGAAGTT AGCTTTGGCC TATTCATAAT	GGGAAAGCGT GTCGGATTAT GATAAGTTCTA GAAAAAGTTA AGAAACTTGT GTTACCGAAC CCCAAAACAG TCCTTTTTTC	TACTAGCTGG CGGCAAGAAT AAGTGCGTGT AAAGTGGCTT TCGCTAACGT TCGTTATTGA ATCGTGGCAA GGCTGGGCTC TATCGGACCT TTTGCTCATTG
66 66 66 66 66 66	5301 5361 5421 5481 5541 6661 6721 6781	TGGATATTTA GCATGCTGTA GAAAAGTAAA CATTCCTGTT ATTTAGCTAC TATTTTCTCT CAGAAGATTC CTCGGCAGGT GGAAAATAAT	ATGTCAGAGT GTTTTAGATA GCCCATCCTT CAATTTAAAC TCGATAGGCG TTTAAACCTA GATGATGCCT AGTTTCATT TTAGTCACAC GTAGACAAAT GGTAAAATTC	TGAGAGCCTT CAAAATATGA ACGGTTCATT ATGGGGAGTG GATCTAAGAA ACATGGACGA CGCACAGAAT TTTTAACTAC ATAGAGAAGG GAAGCGGGTT	CAGTGCCCA AGCATTTGTA GCAAGGGTCT GCTGTACCAT CCCTTTCATT CTACTGCAAT GGAGGATAGC TGAAGAAGTT AGCTTTGGCC TATTCATAAT TGATGTAGCC	GGGAAAGCGT GTCGGATTAT GATAAGTTCTA GAAAAAGTTA AGAAACTTGT CCCAAAACAG TCCTTTTTTG TTAGCACAAG GCGGCAGCAT	TACTAGCTGG CGGCAAGAAT AAGTGCGTGT AAAGTGGCTT TCGCTAACGT TCGTTATTGA ATCGTGGCAA GGCTGGGCTC TATCGGACCT TTTGCTCATTG ATGGATCTAT
66 66 66 66 66 66	5301 5361 5421 5481 5541 6661 6721 6781	TGGATATTTA GCATGCTGTA GAAAAGTAAA CATTCCTGTT ATTTAGCTAC TATTTTCTCT CAGAAGATTC CTCGGCAGGT GGAAAATAAT	ATGTCAGAGT GTTTTAGATA GCCCATCCTT CAATTTAAAC TCGATAGGCG TTTAAACCTA GATGATGCCT AGTTTCATT TTAGTCACAC GTAGACAAAT GGTAAAATTC	TGAGAGCCTT CAAAATATGA ACGGTTCATT ATGGGGAGTG GATCTAAGAA ACATGGACGA CGCACAGAAT TTTTAACTAC ATAGAGAAGG GAAGCGGGTT	CAGTGCCCA AGCATTTGTA GCAAGGGTCT GCTGTACCAT CCCTTTCATT CTACTGCAAT GGAGGATAGC TGAAGAAGTT AGCTTTGGCC TATTCATAAT TGATGTAGCC	GGGAAAGCGT GTCGGATTAT GATAAGTTCTA GAAAAAGTTA AGAAACTTGT CCCAAAACAG TCCTTTTTTG TTAGCACAAG GCGGCAGCAT	TACTAGCTGG CGGCAAGAAT AAGTGCGTGT AAAGTGGCTT TCGCTAACGT TCGTTATTGA ATCGTGGCAA GGCTGGGCTC TATCGGACCT TTTGCTCATTG

6961 TTACGGCAGT AAACTGGCGC ATTTGGTTGA TGAAGAAGAC TGGAATATTA CGATTAAAAG

	-43-							
7021	TAACCATTTA	CCTTCGGGAT	TAACTTTATG	GATGGGCGAT	ATTAAGAATG	GTTCAGAAAC		
7081	AGTAAAACTG	GTCCAGAAGG	TAAAAAATTG	GTATGATTCG	CATATGCCAG	AAAGCTTGAA		
7141	AATATATACA	GAACTCGATC	ATGCAAATTC	TAGATTTATG	GATGGACTAT	CTAAACTAGA		
7201	TCGCTTACAC	GAGACTCATG	ACGATTACAG	CGATCAGATA	TTTGAGTCTC	TTGAGAGGAA		
7261	TGACTGTACC	TGTCAAAAGT	ATCCTGAAAT	CACAGAAGTT	AGAGATGCAG	TTGCCACAAT		
7321	TAGACGTTCC	TTTAGAAAAA	TAACTAAAGA	ATCTGGTGCC	GATATCGAAC	CTCCCGTACA		
7381	AACTAGCTTA	TTGGATGATT	GCCAGACCTT	AAAAGGAGTT	CTTACTTGCT	TAATACCTGG		
7441	TGCTGGTGGT	TATGACGCCA	TTGCAGTGAT	TACTAAGCAA	GATGTTGATC	TTAGGGCTCA		
7501	AACCGCTAAT	GACAAAAGAT	TTTCTAAGGT	TCAATGGCTG	GATGTAACTC	AGGCTGACTG		
		Stop	Phosphomeva	lonate Kinas	se .I	 		
7561	GGGTGTTAGG			TTATCTTGAT	-	TAATACTCAT		
,501	00010111100	111101111111			RBS			
	—▶ Start 1	Mevalonate	Pyrophospha	te Decarbox	ylase			
7621	-			CGTCAACATC		AGTATTGGGG		
7681				CAATTCGTCC				

G 7741 AGATGACCTC AGAACGTTGA CCTCTGCGGC TACTGCACCT GAGTTTGAAC GCGACACTTT 7801 GTGGTTAAAT GGAGAACCAC ACAGCATCGA CAATGAAAGA ACTCAAAATT GTCTGCGCGA 7861 CCTACGCCAA TTAAGAAAGG AAATGGAATC GAAGGACGCC TCATTGCCCA CATTATCTCA 7921 ATGGAAACTC CACATTGTCT CCGAAAATAA CTTTCCTACA GCAGCTGGTT TAGCTTCCTC 7981 CGCTGCTGGC TTTGCTGCAT TGGTCTCTGC AATTGCTAAG TTATACCAAT TACCACAGTC 8041 AACTTCAGAA ATATCTAGAA TAGCAAGAAA GGGGTCTGGT TCAGCTTGTA GATCGTTGTT 8101 TGGCGGATAC GTGGCCTGGG AAATGGGAAA AGCTGAAGAT GGTCATGATT CCATGGCAGT 8161 ACAAATCGCA GACAGCTCTG ACTGGCCTCA GATGAAAGCT TGTGTCCTAG TTGTCAGCGA 8221 TATTAAAAAG GATGTGAGTT CCACTCAGGG TATGCAATTG ACCGTGGCAA CCTCCGAACT 8281 ATTTAAAGAA AGAATTGAAC ATGTCGTACC AAAGAGATTT GAAGTCATGC GTAAAGCCAT 8341 TGTTGAAAAA GATTTCGCCA CCTTTGCAAA GGAAACAATG ATGGATTCCA ACTCTTTCCA 8401 TGCCACATGT TTGGACTCTT TCCCTCCAAT ATTCTACATG AATGACACTT CCAAGCGTAT 8461 CATCAGTTGG TGCCACACCA TTAATCAGTT TTACGGAGAA ACAATCGTTG CATACACGTT 8521 TGATGCAGGT CCAAATGCTG TGTTGTACTA CTTAGCTGAA AATGAGTCGA AACTCTTTGC 8581 ATTTATCTAT AAATTGTTTG GCTCTGTTCC TGGATGGGAC AAGAAATTTA CTACTGAGCA 8641 GCTTGAGGCT TTCAACCATC AATTTGAATC ATCTAACTTT ACTGCACGTG AATTGGATCT 8701 TGAGTTGCAA AAGGATGTTG CCAGAGTGAT TTTAACTCAA GTCGGTTCAG GCCCACAAGA

Stop Mevalonate Pyrophosphate Decarboxylase —— 8761 AACAAACGAA TCTTTGATTG ACGCAAAGAC TGGTCTACCA AAGGAATAAC TGCAGGCATG

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		-	Strong ri	bosomal rrn	B terminato	rs	
	8821	CAAGCTTGGC '	TGTTTTGGCG	GATGAGAGAA	GATTTTCAGC	CTGATACAGA	TTAAATCAGA
	8881	ACGCAGAAGC	GGTCTGATAA	AACAGAATTT	GCCTGGCGGC	AGTAGCGCGG	TGGTCCCACC
5	8941	TGACCCCATG	CCGAACTCAG	AAGTGAAACG	CCGTAGCGCC	GATGGTAGTG	TGGGGTCTCC
	9001	CCATGCGAGA	GTAGGGAACT	GCCAGGCATC	AAATAAAACG	AAAGGCTCAG	TCGAAAGACT
	9061	GGGCCTTTCG	TTTTATCTGT	TGTTTGTCGG	TGAACGCTCT	CCTGAGTAGG	ACAAATCCGC
	9121	CGGGAGCGGA	TTTGAACGTT	GCGAAGCAAC	GGCCCGGAGG	GTGGCGGGCA	GGACGCCCGC
	9181	CATAAACTGC	CAGGCATCAA	ATTAAGCAGA	AGGCCATCCT	GACGGATGGC	CTTTTTGCGT
10	9241	TTCTACAAAC	TCT				
7 7 7 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	SEQ ID NO) 8 "M	EVT" operon	nucleotide sec	quence		
- Contraction (P _{BAD} Promoter					
15 <u>—</u>	1	GACGCTTTTT	ATCGCAACTC	TCTACTGTTT	CTCCATACCC	GTTTTTTTGG	GCTAGCAGGA
A Parameter Communication of the Communication of t							
Antiqued a control of the control of						RBS	
Franchisco Garage Control of the Con	61	GGAATTCACC	ATGGTACCCG	GGGATCCTCT	AGAGTCGACT	AGGAGGAATA	TAAAATGAAA
20	Start	Acetoacetyl-C	OA thiolase				
Application of the control of the co	121	AATTGTGTCA	TCGTCAGTGC	GGTACGTACT	GCTATCGGTA	GTTTTAACGG	TTCACTCGCT
Applications of the control of the c							
	181	TCCACCAGCG	CCATCGACCT	GGGGGCGACA	GTAATTAAAG	CCGCCATTGA	ACGTGCAAAA
-	241	ATCGATTCAC	AACACGTTGA	TGAAGTGATT	ATGGGTAACG	TGTTACAAGC	CGGGCTGGGG
25	301	CAAAATCCGG	CGCGTCAGGC	ACTGTTAAAA	AGCGGGCTGG	CAGAAACGGT	GTGCGGATTC
		ACGGTCAATA					
	421	CAGGCAGGTC	AGGCGCAGAG	CATTGTGGCG	GGGGGTATGG	AAAATATGAG	TTTAGCCCCC
		TACTTACTCG					
		GTAATCCTGC					
30		GAAAACGTGG					
		TCACAGCGTA					
		GTAAATGTTG					
	701	3 3 mm (2 3 3 C/C/C	CTGAAGCGTT	' AGGTGCATTG	CGCCCGGCCT	TCGATAAAGC	AGGAACAGTC
	841	ACCGCTGGGA	ACGCGTCTGG				
35	841 901	ACCGCTGGGA TCTGCGGCGC	ACGCGTCTGG TGGCAGCAGG	CCTTACCCC	CTGGCTCGCA	. TTAAAAGTTA	

1021	CTGGCGGGGC	${\tt TGCAACTGGC}$	${\tt GGATATTGAT}$	CTCATTGAGG	CTAATGAAGC	ATTTGCTGCA
1081	CAGTTCCTTG	CCGTTGGGAA	AAACCTGGGC	TTTGATTCTG	AGAAAGTGAA	TGTCAACGGC
1141	GGGGCCATCG	CGCTCGGGCA	TCCTATCGGT	GCCAGTGGTG	CTCGTATTCT	GGTCACACTA
1201	TTACATGCCA	TGCAGGCACG	CGATAAAACG	CTGGGGCTGG	CAACACTGTG	CATTGGCGGC

Stop Acetoacetyl-CoA Thiolase Start

1261 GGTCAGGGAA TTGCGATGGT GATTGAACGG TTGAATTAAG GAGGACAGCT AAATGAAACT

RBS

HMG-CoA Synthase 1321 CTCAACTAAA CTTTGTTGGT GTGGTATTAA AGGAAGACTT AGGCCGCAAA AGCAACAACA 1381 ATTACACAAT ACAAACTTGC AAATGACTGA ACTAAAAAAA CAAAAGACCG CTGAACAAAA 1441 AACCAGACCT CAAAATGTCG GTATTAAAGG TATCCAAATT TACATCCCAA CTCAATGTGT 1501 CAACCAATCT GAGCTAGAGA AATTTGATGG CGTTTCTCAA GGTAAATACA CAATTGGTCT 1561 GGGCCAAACC AACATGTCTT TTGTCAATGA CAGAGAAGAT ATCTACTCGA TGTCCCTAAC 1621 TGTTTTGTCT AAGTTGATCA AGAGTTACAA CATCGACACC AACAAAATTG GTAGATTAGA 1681 AGTCGGTACT GAAACTCTGA TTGACAAGTC CAAGTCTGTC AAGTCTGTCT TGATGCAATT 1741 GTTTGGTGAA AACACTGACG TCGAAGGTAT TGACACGCTT AATGCCTGTT ACGGTGGTAC 1801 CAACGCGTTG TTCAACTCTT TGAACTGGAT TGAATCTAAC GCATGGGATG GTAGAGACGC 1861 CATTGTAGTT TGCGGTGATA TTGCCATCTA CGATAAGGGT GCCGCAAGAC CAACCGGTGG 1921 TGCCGGTACT GTTGCTATGT GGATCGGTCC TGATGCTCCA ATTGTATTTG ACTCTGTAAG 1981 AGCTTCTTAC ATGGAACACG CCTACGATTT TTACAAGCCA GATTTCACCA GCGAATATCC 2041 TTACGTCGAT GGTCATTTTT CATTAACTTG TTACGTCAAG GCTCTTGATC AAGTTTACAA 2101 GAGTTATTCC AAGAAGGCTA TTTCTAAAGG GTTGGTTAGC GATCCCGCTG GTTCGGATGC 2161 TTTGAACGTT TTGAAATATT TCGACTACAA CGTTTTCCAT GTTCCAACCT GTAAATTGGT 2221 CACAAAATCA TACGGTAGAT TACTATATAA CGATTTCAGA GCCAATCCTC AATTGTTCCC 2281 AGAAGTTGAC GCCGAATTAG CTACTCGCGA TTATGACGAA TCTTTAACCG ATAAGAACAT 2341 TGAAAAACT TTTGTTAATG TTGCTAAGCC ATTCCACAAA GAGAGAGTTG CCCAATCTTT 2401 GATTGTTCCA ACAAACACAG GTAACATGTA CACCGCATCT GTTTATGCCG CCTTTGCATC 2461 TCTATTAAAC TATGTTGGAT CTGACGACTT ACAAGGCAAG CGTGTTGGTT TATTTTCTTA 2521 CGGTTCCGGT TTAGCTGCAT CTCTATATTC TTGCAAAATT GTTGGTGACG TCCAACATAT 2581 TATCAAGGAA TTAGATATTA CTAACAAATT AGCCAAGAGA ATCACCGAAA CTCCAAAGGA 2641 TTACGAAGCT GCCATCGAAT TGAGAGAAAA TGCCCATTTG AAGAAGAACT TCAAACCTCA 2701 AGGTTCCATT GAGCATTTGC AAAGTGGTGT TTACTACTTG ACCAACATCG ATGACAAATT

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	Stop HMG-CoA synthase —— Start Truncate	d
	2761 TAGAAGATCT TACGATGTTA AAAAATAAGG AGGATTACAC TATGGTTTTA ACCAATAAAA	
	RBS	
	HMG-CoA reductase	
5	2821 CAGTCATTTC TGGATCGAAA GTCAAAAGTT TATCATCTGC GCAATCGAGC TCATCAGGAC	
	2881 CTTCATCATC TAGTGAGGAA GATGATTCCC GCGATATTGA AAGCTTGGAT AAGAAAATAC	
	2941 GTCCTTTAGA AGAATTAGAA GCATTATTAA GTAGTGGAAA TACAAAACAA TTGAAGAACA	
	3001 AAGAGGTCGC TGCCTTGGTT ATTCACGGTA AGTTACCTTT GTACGCTTTG GAGAAAAAAT	
	3061 TAGGTGATAC TACGAGAGCG GTTGCGGTAC GTAGGAAGGC TCTTTCAATT TTGGCAGAAG	
10	3121 CTCCTGTATT AGCATCTGAT CGTTTACCAT ATAAAAATTA TGACTACGAC CGCGTATTTG	
	3181 GCGCTTGTTG TGAAAATGTT ATAGGTTACA TGCCTTTGCC CGTTGGTGTT ATAGGCCCCT	
2 a	3241 TGGTTATCGA TGGTACATCT TATCATATAC CAATGGCAAC TACAGAGGGT TGTTTGGTAG	
POT MANAGEMENT OF THE PARTY OF	3301 CTTCTGCCAT GCGTGGCTGT AAGGCAATCA ATGCTGGCGG TGGTGCAACA ACTGTTTTAA	
	3361 CTAAGGATGG TATGACAAGA GGCCCAGTAG TCCGTTTCCC AACTTTGAAA AGATCTGGTG	
15	3421 CCTGTAAGAT ATGGTTAGAC TCAGAAGAGG GACAAAACGC AATTAAAAAA GCTTTTAACT	
Si al di Si	3481 CTACATCAAG ATTTGCACGT CTGCAACATA TTCAAACTTG TCTAGCAGGA GATTTACTCT	
Controlled	3541 TCATGAGATT TAGAACAACT ACTGGTGACG CAATGGGTAT GAATATGATT TCTAAAGGTG	
	3601 TCGAATACTC ATTAAAGCAA ATGGTAGAAG AGTATGGCTG GGAAGATATG GAGGTTGTCT	
OF THE STATE OF TH	3661 CCGTTTCTGG TAACTACTGT ACCGACAAAA AACCAGCTGC CATCAACTGG ATCGAAGGTC	
2014	3721 GTGGTAAGAG TGTCGTCGCA GAAGCTACTA TTCCTGGTGA TGTTGTCAGA AAAGTGTTAA	
A STATEMENT OF THE STAT	3781 AAAGTGATGT TTCCGCATTG GTTGAGTTGA ACATTGCTAA GAATTTGGTT GGATCTGCAA	
	3841 TGGCTGGGTC TGTTGGTGGA TTTAACGCAC ATGCAGCTAA TTTAGTGACA GCTGTTTTCT	
	3901 TGGCATTAGG ACAAGATCCT GCACAAAATG TTGAAAGTTC CAACTGTATA ACATTGATGA	
	3961 AAGAAGTGGA CGGTGATTTG AGAATTTCCG TATCCATGCC ATCCATCGAA GTAGGTACCA	
25	4021 TCGGTGGTGG TACTGTTCTA GAACCACAAG GTGCCATGTT GGACTTATTA GGTGTAAGAG	
	4081 GCCCGCATGC TACCGCTCCT GGTACCAACG CACGTCAATT AGCAAGAATA GTTGCCTGTG	
	4141 CCGTCTTGGC AGGTGAATTA TCCTTATGTG CTGCCCTAGC AGCCGGCCAT TTGGTTCAAA	
	4201 GTCATATGAC CCACAACAGG AAACCTGCTG AACCAACAAA ACCTAACAAT TTGGACGCCA	
30	Stop Truncated HMG-CoA Reductase	
	4261 CTGATATAAA TCGTTTGAAA GATGGGTCCG TCACCTGCAT TAAATCCTAA GTCGACCTGC	
	Strong ribosomal rrnB terminators	

4321 AGGCATGCAA GCTTGGCTGT TTTGGCGGAT GAGAGAAGAT TTTCAGCCTG ATACAGATTA

4381 AATCAGAACG CAGAAGCGGT CTGATAAAAC AGAATTTGCC TGGCGGCAGT AGCGCGGTGG

35

4441 TCCCACCTGA CCCCATGCCG AACTCAGAAG TGAAACGCCG TAGCGCCGAT GGTAGTGTGG 4501 GGTCTCCCCA TGCGAGAGTA GGGAACTGCC AGGCATCAAA TAAAACGAAA GGCTCAGTCG 4561 AAAGACTGGG CCTTTCGTTT TATCTGTTGT TTGTCGGTGA ACGCTCTCCT GAGTAGGACA 4621 AATCCGCCGG GAGCGGATTT GAACGTTGCG AAGCAACGGC CCGGAGGGTG GCGGGCAGGA 4681 CGCCCGCCAT AAACTGCCAG GCATCAAATT AAGCAGAAGG CCATCCTGAC GGATGGCCTT 4741 TTTGCGTTTC TACAAACTCT

"MEVB" operon nucleotide sequence SEQ ID NO 9

Lac Promoter 10 1 GCGCAACGCA ATTAATGTGA GTTAGCTCAC TCATTAGGCA CCCCAGGCTT TACACTTTAT 61 GCTTCCGGCT CGTATGTTGT GTGGAATTGT GAGCGGATAA CAATTTCACA CAGGAAACAG 121 CTATGACCAT GATTACGCCA AGCGCGCAAT TAACCCTCAC TAAAGGGAAC AAAAGCTGGG 181 TACCGGGCCC CCCCTCGAGG TCGACGGTAT CGATAAGCTT GATATCGAAT TCCTGCAGTA ▶ Start Mevalonate Kinase RBS 241 GGAGGAATTA ACCATGTCAT TACCGTTCTT AACTTCTGCA CCGGGAAAGG TTATTATTTT 301 TGGTGAACAC TCTGCTGTGT ACAACAAGCC TGCCGTCGCT GCTAGTGTGT CTGCGTTGAG 361 AACCTACCTG CTAATAAGCG AGTCATCTGC ACCAGATACT ATTGAATTGG ACTTCCCGGA 421 CATTAGCTTT AATCATAAGT GGTCCATCAA TGATTTCAAT GCCATCACCG AGGATCAAGT 481 AAACTCCCAA AAATTGGCCA AGGCTCAACA AGCCACCGAT GGCTTGTCTC AGGAACTCGT 25 541 TAGTCTTTTG GATCCGTTGT TAGCTCAACT ATCCGAATCC TTCCACTACC ATGCAGCGTT 601 TTGTTTCCTG TATATGTTTG TTTGCCTATG CCCCCATGCC AAGAATATTA AGTTTTCTTT 661 AAAGTCTACT TTACCCATCG GTGCTGGGTT GGGCTCAAGC GCCTCTATTT CTGTATCACT 721 GGCCTTAGCT ATGGCCTACT TGGGGGGGTT AATAGGATCT AATGACTTGG AAAAGCTGTC 781 AGAAAACGAT AAGCATATAG TGAATCAATG GGCCTTCATA GGTGAAAAGT GTATTCACGG 30 841 TACCCCTTCA GGAATAGATA ACGCTGTGGC CACTTATGGT AATGCCCTGC TATTTGAAAA 901 AGACTCACAT AATGGAACAA TAAACACAAA CAATTTTAAG TTCTTAGATG ATTTCCCAGC 961 CATTCCAATG ATCCTAACCT ATACTAGAAT TCCAAGGTCT ACAAAAGATC TTGTTGCTCG 1021 CGTTCGTGTG TTGGTCACCG AGAAATTTCC TGAAGTTATG AAGCCAATTC TAGATGCCAT 1081 GGGTGAATGT GCCCTACAAG GCTTAGAGAT CATGACTAAG TTAAGTAAAT GTAAAGGCAC

1141 CGATGACGAG GCTGTAGAAA CTAATAATGA ACTGTATGAA CAACTATTGG AATTGATAAG

	1201 AATAAATCAT GGACTGCTTG TCTCAATCGG TGTTTCTCAT CCTGGATTAG AACTTAT	AATT
	1261 AAATCTGAGC GATGATTTGA GAATTGGCTC CACAAAACTT ACCGGTGCTG GTGGCGG	3CGG
	1321 TTGCTCTTTG ACTTTGTTAC GAAGAGACAT TACTCAAGAG CAAATTGACA GCTTCAF	AAAA
	1381 GAAATTGCAA GATGATTTTA GTTACGAGAC ATTTGAAACA GACTTGGGTG GGACTGG	JCTG
5	1441 CTGTTTGTTA AGCGCAAAAA ATTTGAATAA AGATCTTAAA ATCAAATCCC TAGTATT	rcca
	1501 ATTATTTGAA AATAAAACTA CCACAAAGCA ACAAATTGAC GATCTATTAT TGCCAGC	GAAA
	Stop Mevalonate Kinase Start Phosphomeval	lonate
	1561 CACGAATTTA CCATGGACTT CATAGGAGGC AGATCAAATG TCAGAGTTGA GAGCCTT	FCAG
10	RBS	
	Kinase	
	1621 TGCCCCAGGG AAAGCGTTAC TAGCTGGTGG ATATTTAGTT TTAGATACAA AATATGA	AAGC
or an	1681 ATTTGTAGTC GGATTATCGG CAAGAATGCA TGCTGTAGCC CATCCTTACG GTTCAT	IGCA
	1741 AGGGTCTGAT AAGTTTGAAG TGCGTGTGAA AAGTAAACAA TTTAAAGATG GGGAGT	GGCT
15	1801 GTACCATATA AGTCCTAAAA GTGGCTTCAT TCCTGTTTCG ATAGGCGGAT CTAAGA	ACCC
Controlled	1861 TTTCATTGAA AAAGTTATCG CTAACGTATT TAGCTACTTT AAACCTAACA TGGACG	ACTA
	1921 CTGCAATAGA AACTTGTTCG TTATTGATAT TTTCTCTGAT GATGCCTACC ATTCTC	AGGA
	1981 GGATAGCGTT ACCGAACATC GTGGCAACAG AAGATTGAGT TTTCATTCGC ACAGAA	TTGA
Table of the second sec	2041 AGAAGTTCCC AAAACAGGGC TGGGCTCCTC GGCAGGTTTA GTCACAGTTT TAACTA	CAGC
20	2101 TTTGGCCTCC TTTTTTGTAT CGGACCTGGA AAATAATGTA GACAAATATA GAGAAG	
	2161 TCATAATTTA GCACAAGTTG CTCATTGTCA AGCTCAGGGT AAAATTGGAA GCGGGT	
Parliment Control of the Control of	2221 TGTAGCGGCG GCAGCATATG GATCTATCAG ATATAGAAGA TTCCCACCCG CATTAA	
CONTRACTOR OF THE CONTRACTOR O	2281 TAATTTGCCA GATATTGGAA GTGCTACTTA CGGCAGTAAA CTGGCGCATT TGGTTG	
	2341 AGAAGACTGG AATATTACGA TTAAAAGTAA CCATTTACCT TCGGGATTAA CTTTAT	
25	2401 GGGCGATATT AAGAATGGTT CAGAAACAGT AAAACTGGTC CAGAAGGTAA AAAATT	
	2461 TGATTCGCAT ATGCCAGAAA GCTTGAAAAT ATATACAGAA CTCGATCATG CAAATT	
	2521 ATTTATGGAT GGACTATCTA AACTAGATCG CTTACACGAG ACTCATGACG ATTACA	
	2581 TCAGATATTT GAGTCTCTTG AGAGGAATGA CTGTACCTGT CAAAAGTATC CTGAAA	
	2641 AGAAGTTAGA GATGCAGTTG CCACAATTAG ACGTTCCTTT AGAAAAATAA CTAAAG	
30	2701 TGGTGCCGAT ATCGAACCTC CCGTACAAAC TAGCTTATTG GATGATTGCC AGACCT	
	2761 AGGAGTTCTT ACTTGCTTAA TACCTGGTGC TGGTGGTTAT GACGCCATTG CAGTGA	
	2821 TAAGCAAGAT GTTGATCTTA GGGCTCAAAC CGCTAATGAC AAAAGATTTT CTAAGG	;TTCA
	Stop Phosphomeval	
35	2881 ATGGCTGGAT GTAACTCAGG CTGACTGGGG TGTTAGGAAA GAAAAAGATC CGGAAA	CTTA

10

20 4

25

30

35

Kinase Start Mevalonate Pyrophosphate

2941 TCTTGATAAA TAGGAGGTAA TACTCATGAC CGTTTACACA GCATCCGTTA CCGCACCCGT

RBS

Decarboxylase 3001 CAACATCGCA ACCCTTAAGT ATTGGGGGAA AAGGGACACG AAGTTGAATC TGCCCACCAA 3061 TTCGTCCATA TCAGTGACTT TATCGCAAGA TGACCTCAGA ACGTTGACCT CTGCGGCTAC 3121 TGCACCTGAG TTTGAACGCG ACACTTTGTG GTTAAATGGA GAACCACACA GCATCGACAA 3181 TGAAAGAACT CAAAATTGTC TGCGCGACCT ACGCCAATTA AGAAAGGAAA TGGAATCGAA 3241 GGACGCCTCA TTGCCCACAT TATCTCAATG GAAACTCCAC ATTGTCTCCG AAAATAACTT 3301 TCCTACAGCA GCTGGTTTAG CTTCCTCCGC TGCTGGCTTT GCTGCATTGG TCTCTGCAAT 3361 TGCTAAGTTA TACCAATTAC CACAGTCAAC TTCAGAAATA TCTAGAATAG CAAGAAAGGG 3421 GTCTGGTTCA GCTTGTAGAT CGTTGTTTGG CGGATACGTG GCCTGGGAAA TGGGAAAAGC 3481 TGAAGATGGT CATGATTCCA TGGCAGTACA AATCGCAGAC AGCTCTGACT GGCCTCAGAT 3541 GAAAGCTTGT GTCCTAGTTG TCAGCGATAT TAAAAAGGAT GTGAGTTCCA CTCAGGGTAT 3601 GCAATTGACC GTGGCAACCT CCGAACTATT TAAAGAAAGA ATTGAACATG TCGTACCAAA 3661 GAGATTTGAA GTCATGCGTA AAGCCATTGT TGAAAAAGAT TTCGCCACCT TTGCAAAGGA 3721 AACAATGATG GATTCCAACT CTTTCCATGC CACATGTTTG GACTCTTTCC CTCCAATATT 3781 CTACATGAAT GACACTTCCA AGCGTATCAT CAGTTGGTGC CACACCATTA ATCAGTTTTA 3841 CGGAGAAACA ATCGTTGCAT ACACGTTTGA TGCAGGTCCA AATGCTGTGT TGTACTACTT 3901 AGCTGAAAAT GAGTCGAAAC TCTTTGCATT TATCTATAAA TTGTTTGGCT CTGTTCCTGG 3961 ATGGGACAAG AAATTTACTA CTGAGCAGCT TGAGGCTTTC AACCATCAAT TTGAATCATC

Stop Mevalonate Pyrophosphate
4081 AACTCAAGTC GGTTCAGGCC CACAAGAAAC AAACGAATCT TTGATTGACG CAAAGACTGG

4021 TAACTTTACT GCACGTGAAT TGGATCTTGA GTTGCAAAAG GATGTTGCCA GAGTGATTTT

Decarboxylase -

4141TCTACCAAAGGAATAACTGCAGCCCGGGGGATCCACTAGTTCTAGAGCGGCCGCCACCGC4201GGTGGAGCTCCAATTCGCCCTATAGTGAGTCGTATTACGCGCGCTCACTGGCCGTCGTTT4261TACAACGTCGTGACTGGGAAAACCCTGGCGTTACCCAACTTAATCGCCTTGCAGCACATC4321CCCCTTTCGCCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCGCCTTCCCAACAGT4381TGCGCAGCCTGAATGGCGAATGGAAATTGTAAGCGTTAATATTTTGTTAAAATTCGCGTT4441AAATTTTTGTTAAATCAGCTCATTTTTAACCAATAGGCCGA

SEQ ID NO 10 Isopentyl pyrophosphate isomerase (idi) nucleotide sequence

1 ATGCAAACGG AACACGTCAT TTTATTGAAT GCACAGGGAG TTCCCACGGG TACGCTGGAA

61	AAGTATGCCG	CACACACGGC	AGACACCCGC	TTACATCTCG	CGTTCTCCAG	TTGGCTGTTT
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181	GTGTGGACTA	ACTCGGTTTG	TGGGCACCCA	CAACTGGGAG	AAAGCAACGA	AGACGCAGTG
241	ATCCGCCGTT	GCCGTTATGA	GCTTGGCGTG	GAAATTACGC	CTCCTGAATC	TATCTATCCT
301	GACTTTCGCT	ACCGCGCCAC	CGATCCGAGT	GGCATTGTGG	AAAATGAAGT	GTGTCCGGTA
361	TTTGCCGCAC	GCACCACTAG	TGCGTTACAG	ATCAATGATG	ATGAAGTGAT	GGATTATCAA
421	TGGTGTGATT	TAGCAGATGT	ATTACACGGT	ATTGATGCCA	CGCCGTGGGC	GTTCAGTCCG
481	TGGATGGTGA	TGCAGGCGAC	AAATCGCGAA	GCCAGAAAAC	GATTATCTGC	ATTTACCCAG
541	CTTAAATAA					
SEQ ID NO	11 Fai	rnesyl pyropho	osphate syntha	ase (ispA) nuc	leotide sequer	nce
•				GTTAAGCAGG		
61	TTTATCGCCC	CACTGCCCTT	TCAGAACACT	CCCGTGGTCG	AAACCATGCA	GTATGGCGCA
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241	TTAATTCATG	ATGATTTACC	GGCAATGGAT	GATGACGATC	TGCGTCGCGG	TTTGCCAACC
301	TGCCATGTGA	AGTTTGGCGA	AGCAAACGCG	ATTCTCGCTG	GCGACGCTTT	ACAAACGCTG
361	GCGTTCTCGA	TTTTAAGCGA	TGCCGATATG	CCGGAAGTGT	CGGACCGCGA	CAGAATTTCG
421	ATGATTTCTG	AACTGGCGAG	CGCCAGTGGT	ATTGCCGGAA	TGTGCGGTGG	TCAGGCATTA
481	GATTTAGACG	CGGAAGGCAA	ACACGTACCT	CTGGACGCGC	TTGAGCGTAT	TCATCGTCAT
541	AAAACCGGCG	CATTGATTCG	CGCCGCCGTT	CGCCTTGGTG	CATTAAGCGC	CGGAGATAAA
601	GGACGTCGTG	CTCTGCCGGT	ACTCGACAAG	TATGCAGAGA	GCATCGGCCT	TGCCTTCCAG
661	GTTCAGGATG	ACATCCTGGA	TGTGGTGGGA	GATACTGCAA	CGTTGGGAAA	ACGCCAGGGT
721	GCCGACCAGC	: AACTTGGTAA	AAGTACCTAC	CCTGCACTTC	TGGGTCTTGA	GCAAGCCCGG
781	AAGAAAGCCC	GGGATCTGAT	CGACGATGCC	CGTCAGTCGC	TGAAACAACT	GGCTGAACAG
841	TCACTCGATA	CCTCGGCACT	GGAAGCGCTA	GCGGACTACA	TCATCCAGCG	AATAAATAAT
SEQ ID NO	O 12 "N	/IBI" operon n	ucleotide seq	uence		
	Lac Promot	er				
1	GCGCAACGC	A ATTAATGTGA	GTTAGCTCAC	TCATTAGGCA	CCCCAGGCTT	TACACTTTAT
61	GCTTCCGGCT	r CGTATGTTGT	GTGGAATTGT	GAGCGGATAA	CAATTTCACA	CAGGAAACAG
121	CTATGACCA:	r gattacgcca	A AGCGCGCAA	TAACCCTCAC	TAAAGGGAAC	AAAAGCTGGG

181 TACCGGGCCC CCCCTCGAGG TCGACGGTAT CGATAAGCTT GATATCGAAT TCCTGCAGTA

	181 TACCGGGCCC CCCCTCGAGG ICGACGGIAI CGAIAAGCII GAIAICGAAI ICCIGGAC	J 1
5	RBS Start Mevalonate Kinase	
•	241 GGAGGAATTA ACCATGTCAT TACCGTTCTT AACTTCTGCA CCGGGAAAGG TTATTAT	TTT
	301 TGGTGAACAC TCTGCTGTGT ACAACAAGCC TGCCGTCGCT GCTAGTGTGT CTGCGTTC	
	361 AACCTACCTG CTAATAAGCG AGTCATCTGC ACCAGATACT ATTGAATTGG ACTTCCCC	
10	421 CATTAGCTTT AATCATAAGT GGTCCATCAA TGATTTCAAT GCCATCACCG AGGATCA	
	481 AAACTCCCAA AAATTGGCCA AGGCTCAACA AGCCACCGAT GGCTTGTCTC AGGAACT	
	541 TAGTCTTTTG GATCCGTTGT TAGCTCAACT ATCCGAATCC TTCCACTACC ATGCAGC	
C C C C C C C C C C C C C C C C C C C	601 TTGTTTCCTG TATATGTTTG TTTGCCTATG CCCCCATGCC AAGAATATTA AGTTTTC	TTT
i marantana i mar	661 AAAGTCTACT TTACCCATCG GTGCTGGGTT GGGCTCAAGC GCCTCTATTT CTGTATC	ACT
	721 GGCCTTAGCT ATGGCCTACT TGGGGGGGTT AATAGGATCT AATGACTTGG AAAAGCT	GTC
Account 5	781 AGAAAACGAT AAGCATATAG TGAATCAATG GGCCTTCATA GGTGAAAAGT GTATTCA	.CGG
2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	841 TACCCCTTCA GGAATAGATA ACGCTGTGGC CACTTATGGT AATGCCCTGC TATTTGA	AAA
Comments of the comments of th	901 AGACTCACAT AATGGAACAA TAAACACAAA CAATTTTAAG TTCTTAGATG ATTTCCC	:AGC
Section (961 CATTCCAATG ATCCTAACCT ATACTAGAAT TCCAAGGTCT ACAAAAGATC TTGTTGC	TCG
	1021 CGTTCGTGTG TTGGTCACCG AGAAATTTCC TGAAGTTATG AAGCCAATTC TAGATGC	CAT
	1081 GGGTGAATGT GCCCTACAAG GCTTAGAGAT CATGACTAAG TTAAGTAAAT GTAAAGG	}CAC
	1141 CGATGACGAG GCTGTAGAAA CTAATAATGA ACTGTATGAA CAACTATTGG AATTGAT	ZAAG
A consideration of the constraint of the constra	1201 AATAAATCAT GGACTGCTTG TCTCAATCGG TGTTTCTCAT CCTGGATTAG AACTTAT	CTAA
	1261 AAATCTGAGC GATGATTTGA GAATTGGCTC CACAAAACTT ACCGGTGCTG GTGGCGG	3CGG
25	1321 TTGCTCTTTG ACTTTGTTAC GAAGAGACAT TACTCAAGAG CAAATTGACA GCTTCAA	AAA
	1381 GAAATTGCAA GATGATTTTA GTTACGAGAC ATTTGAAACA GACTTGGGTG GGACTGC	JCTG
	1441 CTGTTTGTTA AGCGCAAAAA ATTTGAATAA AGATCTTAAA ATCAAATCCC TAGTATT	rcca
	1501 ATTATTTGAA AATAAAACTA CCACAAAGCA ACAAATTGAC GATCTATTAT TGCCAGC	GAAA
30	Stop Mevalonate Kinase Start Phosphomeval	lonate
	1561 CACGAATTTA CCATGGACTT CATAGGAGGC AGATCAAATG TCAGAGTTGA GAGCCT	TCAG
	RBS	
	Kinase	
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	1741 AGGGTCTGAT AAGTTTGAAG TGCGTGTGAA AAGTAAACAA TTTAAAGATG GGGAGT	GGCT

	1801	GTACCATATA	AGTCCTAAAA	GTGGCTTCAT	TCCTGTTTCG	ATAGGCGGAT	CTAAGAACCC
:	1861	TTTCATTGAA	AAAGTTATCG	CTAACGTATT	TAGCTACTTT	AAACCTAACA	TGGACGACTA
-	1921	CTGCAATAGA	AACTTGTTCG	TTATTGATAT	TTTCTCTGAT	GATGCCTACC	ATTCTCAGGA
:	1981	GGATAGCGTT	ACCGAACATC	GTGGCAACAG	AAGATTGAGT	TTTCATTCGC	ACAGAATTGA
:	2041	AGAAGTTCCC	AAAACAGGGC	TGGGCTCCTC	GGCAGGTTTA	GTCACAGTTT	TAACTACAGC
:	2101	TTTGGCCTCC	TTTTTTGTAT	CGGACCTGGA	AAATAATGTA	GACAAATATA	GAGAAGTTAT
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:	2281	TAATTTGCCA	GATATTGGAA	GTGCTACTTA	CGGCAGTAAA	CTGGCGCATT	TGGTTGATGA
:	2341	AGAAGACTGG	AATATTACGA	TTAAAAGTAA	CCATTTACCT	TCGGGATTAA	CTTTATGGAT
:	2401	GGGCGATATT	AAGAATGGTT	CAGAAACAGT	AAAACTGGTC	CAGAAGGTAA	AAAATTGGTA
:	2461	TGATTCGCAT	ATGCCAGAAA	GCTTGAAAAT	ATATACAGAA	CTCGATCATG	CAAATTCTAG
:	2521	ATTTATGGAT	GGACTATCTA	AACTAGATCG	CTTACACGAG	ACTCATGACG	ATTACAGCGA
:	2581	TCAGATATTT	GAGTCTCTTG	AGAGGAATGA	CTGTACCTGT	CAAAAGTATC	CTGAAATCAC
:	2641	AGAAGTTAGA	GATGCAGTTG	CCACAATTAG	ACGTTCCTTT	AGAAAAATAA	CTAAAGAATC
	2701	TGGTGCCGAT	ATCGAACCTC	CCGTACAAAC	TAGCTTATTG	GATGATTGCC	AGACCTTAAA
	2761	AGGAGTTCTT	ACTTGCTTAA	TACCTGGTGC	TGGTGGTTAT	GACGCCATTG	CAGTGATTAC
	2821	TAAGCAAGAT	GTTGATCTTA	GGGCTCAAAC	CGCTAATGAC	AAAAGATTTT	CTAAGGTTCA
						Stop Phospl	nomevalonate
	2881	ATGGCTGGAT	GTAACTCAGG	CTGACTGGGG	TGTTAGGAAA	Stop Phospl	
	2881	ATGGCTGGAT	GTAACTCAGG	CTGACTGGGG	TGTTAGGAAA		
		ATGGCTGGAT	GTAACTCAGG				CGGAAACTTA
	K	inase ———	 ▶	 	▶ Start Meva	GAAAAAGATC	CGGAAACTTA
	K	inase ———	 ▶	 	▶ Start Meva	GAAAAAGATC	CGGAAACTTA
	K: 2941	inase ———	TAGGAGGTAA	 	▶ Start Meva	GAAAAAGATC	CGGAAACTTA
De	K: 2941 carbo	inase ————————————————————————————————————	TAGGAGGTAA RBS	TACTCATGAC	► Start Meva	GAAAAAGATC	CGGAAACTTA ophosphate CCGCACCCGT
De	Ki 2941 carbo 3001	inase ————————————————————————————————————	TAGGAGGTAA RBS ACCCTTAAGT	TACTCATGAC ATTGGGGGAA	Start Meva	GAAAAAGATC alonate Pyro	CGGAAACTTA Ophosphate CCGCACCCGT TGCCCACCAA
De	K2941 carbo 3001 3061	inase ————————————————————————————————————	TAGGAGGTAA RBS ACCCTTAAGT TCAGTGACTT	TACTCATGAC ATTGGGGGAA TATCGCAAGA	Start Meva CGTTTACACA AAGGGACACG TGACCTCAGA	GAAAAAGATC alonate Pyro GCATCCGTTA AAGTTGAATC	CGGAAACTTA ophosphate CCGCACCCGT TGCCCACCAA CTGCGGCTAC
De	Ki 2941 carbo 3001 3061 3121	inase TCTTGATAAA	TAGGAGGTAA RBS ACCCTTAAGT TCAGTGACTT TTTGAACGCG	TACTCATGAC ATTGGGGGAA TATCGCAAGA ACACTTTGTG	Start Mevander Mevand	GAAAAAGATC alonate Pyro GCATCCGTTA AAGTTGAATC ACGTTGACCT	CGGAAACTTA Dphosphate CCGCACCGT TGCCCACCAA CTGCGGCTAC GCATCGACAA
De	K: 2941 carbo 3001 3061 3121 3181	inase ————————————————————————————————————	TAGGAGGTAA RBS ACCCTTAAGT TCAGTGACTT TTTGAACGCG CAAAATTGTC	TACTCATGAC ATTGGGGGAA TATCGCAAGA ACACTTTGTG TGCGCGACCT	Start Mevander Medical Medical Mevander Medical Medical Medical Mevander Medical Medical Medical Mevander Medical Medical Medical Medical Medical Medical Medical Mevander Medical M	GAAAAAGATC alonate Pyro GCATCCGTTA AAGTTGAATC ACGTTGACCT GAACCACACA	CGGAAACTTA Dphosphate CCGCACCGT TGCCCACCAA CTGCGGCTAC GCATCGACAA TGGAATCGAA
De	Karbo 2941 carbo 3001 3061 3121 3181 3241	inase TCTTGATAAA DXYlase CAACATCGCA TTCGTCCATA TGCACCTGAG TGAAAGAACT GGACGCCTCA	TAGGAGGTAA RBS ACCCTTAAGT TCAGTGACTT TTTGAACGCG CAAAATTGTC TTGCCCACAT	TACTCATGAC ATTGGGGGAA TATCGCAAGA ACACTTTGTG TGCGCGACCT TATCTCAATG	Start Meval CGTTTACACA AAGGGACACG TGACCTCAGA GTTAAATGGA ACGCCAATTA GAAACTCCAC	GAAAAAGATC alonate Pyro GCATCCGTTA AAGTTGAATC ACGTTGACCT GAACCACACA AGAAAGGAAA	CGGAAACTTA Dphosphate CCGCACCGT TGCCCACCAA CTGCGGCTAC GCATCGACAA TGGAATCGAA AAAATAACTT
De	Karbo 2941 carbo 3001 3061 3121 3181 3241 3301	inase TCTTGATAAA DXYlase CAACATCGCA TTCGTCCATA TGCACCTGAG TGAAAGAACT GGACGCCTCA TCCTACAGCA	TAGGAGGTAA RBS ACCCTTAAGT TCAGTGACTT TTTGAACGCG CAAAATTGTC TTGCCCACAT GCTGGTTTAG	TACTCATGAC ATTGGGGGAA TATCGCAAGA ACACTTTGTG TGCGCGACCT TATCTCAATG CTTCCTCCGC	Start Meval CGTTTACACA AAGGGACACG TGACCTCAGA GTTAAATGGA ACGCCAATTA GAAACTCCAC TGCTGGCTTT	GAAAAAGATC alonate Pyro GCATCCGTTA AAGTTGAATC ACGTTGACCT GAACCACACA AGAAAGGAAA ATTGTCTCCG	CGGAAACTTA Dphosphate CCGCACCCGT TGCCCACCAA CTGCGGCTAC GCATCGACAA TGGAATCGACA AAAATAACTT TCTCTGCAAT
De	Karbo 2941 carbo 3001 3061 3121 3181 3241 3301 3361	inase ————————————————————————————————————	TAGGAGGTAA RBS ACCCTTAAGT TCAGTGACTT TTTGAACGCG CAAAATTGTC TTGCCCACAT GCTGGTTTAG TACCAATTAC	TACTCATGAC ATTGGGGGAA TATCGCAAGA ACACTTTGTG TGCGCGACCT TATCTCAATG CTTCCTCCGC CACAGTCAAC	Start Meval CGTTTACACA AAGGGACACG TGACCTCAGA GTTAAATGGA ACGCCAATTA GAAACTCCAC TGCTGGCTTT TTCAGAAATA	GAAAAAGATC alonate Pyro GCATCCGTTA AAGTTGAATC ACGTTGACCT GAACCACACA AGAAAGGAAA ATTGTCTCCG GCTGCATTGG	CGGAAACTTA Dphosphate CCGCACCGT TGCCCACCAA CTGCGGCTAC GCATCGACAA TGGAATCGAA AAAATAACTT TCTCTGCAAT CAAGAAAGGG
De	Karbo 3001 3061 3121 3181 3241 3301 3361 3421	inase TCTTGATAAA DXYlase CAACATCGCA TTCGTCCATA TGCACCTGAG TGAAAGAACT GGACGCCTCA TCCTACAGCA TGCTACAGCA TGCTACAGCTA	TAGGAGGTAA RBS ACCCTTAAGT TCAGTGACTT TTTGAACGCG CAAAATTGTC TTGCCCACAT GCTGGTTTAG TACCAATTAC GCTTGTAGAT	TACTCATGAC ATTGGGGGAA TATCGCAAGA ACACTTTGTG TGCGCGACCT TATCTCAATG CTTCCTCCGC CACAGTCAAC CGTTGTTTGG	Start Meval CGTTTACACA AAGGGACACG TGACCTCAGA GTTAAATGGA ACGCCAATTA GAAACTCCAC TGCTGGCTTT TTCAGAAATA CGGATACGTG	GAAAAAGATC alonate Pyro GCATCCGTTA AAGTTGAATC ACGTTGACCT GAACCACACA AGAAAGGAAA ATTGTCTCCG GCTGCATTGG TCTAGAATAG	CGGAAACTTA Dphosphate CCGCACCGT TGCCCACCAA CTGCGGCTAC GCATCGACAA TGGAATCGAA AAAATAACTT TCTCTGCAAT CAAGAAAGGG TGGGAAAAGC

3541 GAAAGCTTGT GTCCTAGTTG TCAGCGATAT TAAAAAGGAT GTGAGTTCCA CTCAGGGTAT

-55-
3601 GCAATTGACC GTGGCAACCT CCGAACTATT TAAAGAAAGA ATTGAACATG TCGTACCAAA
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Stop Mevalonate Pyrophosphate
4081 AACTCAAGTC GGTTCAGGCC CACAAGAAAC AAACGAATCT TTGATTGACG CAAAGACTGG
Decarboxylase RBS Start Isopentyl
4141 TCTACCAAAG GAATAACTGC AGCCCGGGAG GAGGATTACT ATATGCAAAC GGAACACGTC
Pyrophosphate Isomerase
4201 ATTTTATTGA ATGCACAGGG AGTTCCCACG GGTACGCTGG AAAAGTATGC CGCACACACG
4261 GCAGACACCC GCTTACATCT CGCGTTCTCC AGTTGGCTGT TTAATGCCAA AGGACAATTA
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4621 GTATTACACG GTATTGATGC CACGCCGTGG GCGTTCAGTC CGTGGATGGT GATGCAGGCG
Stop Isopentyl Pyrophosphate Decarboxylase
4681 ACAAATCGCG AAGCCAGAAA ACGATTATCT GCATTTACCC AGCTTAAATA ACCCGGGGGA
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4981 AGCGTTAATA TTTTGTTAAA ATTCGCGTTA AATTTTTGTT AAATCAGCTC ATTTTTTAAC
5041 CAATAGGCCG A

SEQ ID NO 13 "MBIS" operon nucleotide sequence

Lac Promoter

1 GCGCAACGCA ATTAATGTGA GTTAGCTCAC TCATTAGGCA CCCCAGGCTT TACACTTTAT 5 61 GCTTCCGGCT CGTATGTTGT GTGGAATTGT GAGCGGATAA CAATTTCACA CAGGAAACAG 121 CTATGACCAT GATTACGCCA AGCGCGCAAT TAACCCTCAC TAAAGGGAAC AAAAGCTGGG 10 181 TACCGGGCCC CCCCTCGAGG TCGACGGTAT CGATAAGCTT GATATCGAAT TCCTGCAGTÃ ▶ Start Mevalonate Kinase RBS 241 GGAGGAATTA ACCATGTCAT TACCGTTCTT AACTTCTGCA CCGGGAAAGG TTATTATTTT 301 TGGTGAACAC TCTGCTGTGT ACAACAAGCC TGCCGTCGCT GCTAGTGTGT CTGCGTTGAG 361 AACCTACCTG CTAATAAGCG AGTCATCTGC ACCAGATACT ATTGAATTGG ACTTCCCGGA 421 CATTAGCTTT AATCATAAGT GGTCCATCAA TGATTTCAAT GCCATCACCG AGGATCAAGT 481 AAACTCCCAA AAATTGGCCA AGGCTCAACA AGCCACCGAT GGCTTGTCTC AGGAACTCGT 541 TAGTCTTTTG GATCCGTTGT TAGCTCAACT ATCCGAATCC TTCCACTACC ATGCAGCGTT H 201 601 TTGTTTCCTG TATATGTTTG TTTGCCTATG CCCCCATGCC AAGAATATTA AGTTTTCTTT 661 AAAGTCTACT TTACCCATCG GTGCTGGGTT GGGCTCAAGC GCCTCTATTT CTGTATCACT 721 GGCCTTAGCT ATGGCCTACT TGGGGGGGTT AATAGGATCT AATGACTTGG AAAAGCTGTC 781 AGAAAACGAT AAGCATATAG TGAATCAATG GGCCTTCATA GGTGAAAAGT GTATTCACGG 841 TACCCCTTCA GGAATAGATA ACGCTGTGGC CACTTATGGT AATGCCCTGC TATTTGAAAA 901 AGACTCACAT AATGGAACAA TAAACACAAA CAATTTTAAG TTCTTAGATG ATTTCCCAGC 25 961 CATTCCAATG ATCCTAACCT ATACTAGAAT TCCAAGGTCT ACAAAAGATC TTGTTGCTCG 1021 CGTTCGTGTG TTGGTCACCG AGAAATTTCC TGAAGTTATG AAGCCAATTC TAGATGCCAT 1081 GGGTGAATGT GCCCTACAAG GCTTAGAGAT CATGACTAAG TTAAGTAAAT GTAAAGGCAC 1141 CGATGACGAG GCTGTAGAAA CTAATAATGA ACTGTATGAA CAACTATTGG AATTGATAAG 1201 AATAAATCAT GGACTGCTTG TCTCAATCGG TGTTTCTCAT CCTGGATTAG AACTTATTAA 30 1261 AAATCTGAGC GATGATTTGA GAATTGGCTC CACAAAACTT ACCGGTGCTG GTGGCGGCGG 1321 TTGCTCTTTG ACTTTGTTAC GAAGAGACAT TACTCAAGAG CAAATTGACA GCTTCAAAAA 1381 GAAATTGCAA GATGATTTTA GTTACGAGAC ATTTGAAACA GACTTGGGTG GGACTGGCTG 1441 CTGTTTGTTA AGCGCAAAAA ATTTGAATAA AGATCTTAAA ATCAAATCCC TAGTATTCCA

1501 ATTATTTGAA AATAAAACTA CCACAAAGCA ACAAATTGAC GATCTATTAT TGCCAGGAAA

	Stop Mevalonate Kinase Start Phosphomevalonate
	1561 CACGAATTTA CCATGGACTT CATAGGAGGC AGATCAAATG TCAGAGTTGA GAGCCTTCAG
	RBS
	Kinase
5	1621 TGCCCCAGGG AAAGCGTTAC TAGCTGGTGG ATATTTAGTT TTAGATACAA AATATGAAGC
	1681 ATTTGTAGTC GGATTATCGG CAAGAATGCA TGCTGTAGCC CATCCTTACG GTTCATTGCA
	1741 AGGGTCTGAT AAGTTTGAAG TGCGTGTGAA AAGTAAACAA TTTAAAGATG GGGAGTGGCT
	1801 GTACCATATA AGTCCTAAAA GTGGCTTCAT TCCTGTTTCG ATAGGCGGAT CTAAGAACCC
	1861 TTTCATTGAA AAAGTTATCG CTAACGTATT TAGCTACTTT AAACCTAACA TGGACGACTA
10	1921 CTGCAATAGA AACTTGTTCG TTATTGATAT TTTCTCTGAT GATGCCTACC ATTCTCAGGA
	1981 GGATAGCGTT ACCGAACATC GTGGCAACAG AAGATTGAGT TTTCATTCGC ACAGAATTGA
	2041 AGAAGTTCCC AAAACAGGGC TGGGCTCCTC GGCAGGTTTA GTCACAGTTT TAACTACAGC
10 pp 20 pp 10 pp	2101 TTTGGCCTCC TTTTTTGTAT CGGACCTGGA AAATAATGTA GACAAATATA GAGAAGTTAT
#	2161 TCATAATTTA GCACAAGTTG CTCATTGTCA AGCTCAGGGT AAAATTGGAA GCGGGTTTGA
15	2221 TGTAGCGGCG GCAGCATATG GATCTATCAG ATATAGAAGA TTCCCACCCG CATTAATCTC
Control of the contro	2281 TAATTTGCCA GATATTGGAA GTGCTACTTA CGGCAGTAAA CTGGCGCATT TGGTTGATGA
Control of the contro	2341 AGAAGACTGG AATATTACGA TTAAAAGTAA CCATTTACCT TCGGGATTAA CTTTATGGAT
Annual Community of the	2401 GGGCGATATT AAGAATGGTT CAGAAACAGT AAAACTGGTC CAGAAGGTAA AAAATTGGTA
	2461 TGATTCGCAT ATGCCAGAAA GCTTGAAAAT ATATACAGAA CTCGATCATG CAAATTCTAG
20	2521 ATTTATGGAT GGACTATCTA AACTAGATCG CTTACACGAG ACTCATGACG ATTACAGCGA
Comments Com	2581 TCAGATATTT GAGTCTCTTG AGAGGAATGA CTGTACCTGT CAAAAGTATC CTGAAATCAC
2 2 2 2 2 2 2 2 2	2641 AGAAGTTAGA GATGCAGTTG CCACAATTAG ACGTTCCTTT AGAAAAATAA CTAAAGAATC
Tomorrow and to the control of the c	2701 TGGTGCCGAT ATCGAACCTC CCGTACAAAC TAGCTTATTG GATGATTGCC AGACCTTAAA
*	2761 AGGAGTTCTT ACTTGCTTAA TACCTGGTGC TGGTGGTTAT GACGCCATTG CAGTGATTAC
25	2821 TAAGCAAGAT GTTGATCTTA GGGCTCAAAC CGCTAATGAC AAAAGATTTT CTAAGGTTCA
	Stop Phosphomevalonate
	2881 ATGGCTGGAT GTAACTCAGG CTGACTGGGG TGTTAGGAAA GAAAAAGATC CGGAAACTTA
30	Kinase — Start Mevalonate Pyrophosphate
	2941 TCTTGATAAA TAGGAGGTAA TACTCATGAC CGTTTACACA GCATCCGTTA CCGCACCCGT
	RBS
	Decarboxylase
	3001 CAACATCGCA ACCCTTAAGT ATTGGGGGAA AAGGGACACG AAGTTGAATC TGCCCACCAA
35	3061 TTCGTCCATA TCAGTGACTT TATCGCAAGA TGACCTCAGA ACGTTGACCT CTGCGGCTAC
	3121 TGCACCTGAG TTTGAACGCG ACACTTTGTG GTTAAATGGA GAACCACACA GCATCGACAA

3181	TGAAAGAACT	CAAAATTGTC	TGCGCGACCT	ACGCCAATTA	AGAAAGGAAA	TGGAATCGAA
3241	GGACGCCTCA	TTGCCCACAT	TATCTCAATG	GAAACTCCAC	ATTGTCTCCG	AAAATAACTT
3301	TCCTACAGCA	GCTGGTTTAG	CTTCCTCCGC	TGCTGGCTTT	GCTGCATTGG	TCTCTGCAAT
3361	TGCTAAGTTA	TACCAATTAC	CACAGTCAAC	TTCAGAAATA	TCTAGAATAG	CAAGAAAGGG
3421	GTCTGGTTCA	GCTTGTAGAT	CGTTGTTTGG	CGGATACGTG	GCCTGGGAAA	TGGGAAAAGC
3481	TGAAGATGGT	CATGATTCCA	TGGCAGTACA	AATCGCAGAC	AGCTCTGACT	GGCCTCAGAT
3541	GAAAGCTTGT	GTCCTAGTTG	TCAGCGATAT	TAAAAAGGAT	GTGAGTTCCA	CTCAGGGTAT
3601	GCAATTGACC	GTGGCAACCT	CCGAACTATT	TAAAGAAAGA	ATTGAACATG	TCGTACCAAA
3661	GAGATTTGAA	GTCATGCGTA	AAGCCATTGT	TGAAAAAGAT	TTCGCCACCT	TTGCAAAGGA
3721	AACAATGATG	GATTCCAACT	CTTTCCATGC	CACATGTTTG	GACTCTTTCC	CTCCAATATT
3781	CTACATGAAT	GACACTTCCA	AGCGTATCAT	CAGTTGGTGC	CACACCATTA	ATCAGTTTTA
3841	CGGAGAAACA	ATCGTTGCAT	ACACGTTTGA	TGCAGGTCCA	AATGCTGTGT	TGTACTACTT
3901	AGCTGAAAAT	GAGTCGAAAC	TCTTTGCATT	TATCTATAAA	TTGTTTGGCT	CTGTTCCTGG
3961	ATGGGACAAG	AAATTTACTA	CTGAGCAGCT	TGAGGCTTTC	AACCATCAAT	TTGAATCATC
4021	TAACTTTACT	GCACGTGAAT	TGGATCTTGA	GTTGCAAAAG	GATGTTGCCA	GAGTGATTTT
				Stop I	Mevalonate 1	Pyrophosphate
4081	AACTCAAGTC	GGTTCAGGCC	CACAAGAAAC	-		
4081	AACTCAAGTC	GGTTCAGGCC	CACAAGAAAC	-		
	AACTCAAGTC arboxylase	GGTTCAGGCC		-	TTGATTGACG	
Dec	arboxylase	GGTTCAGGCC GAATAACTGC	I	AAACGAATCT	TTGATTGACG Star	CAAAGACTGG
Dec	arboxylase	>	I	AAACGAATCT	TTGATTGACG Star	CAAAGACTGG
Dec.	arboxylase	GAATAACTGC	I	AAACGAATCT	TTGATTGACG Star	CAAAGACTGG
Dec. 4141 Pyropho	arboxylase TCTACCAAAG sphate Isom	GAATAACTGC	AGCCCGGG <mark>ĀG</mark>	AAACGAATCT RBS	TTGATTGACG ├─► Stan ATATGCAAAC	CAAAGACTGG rt Isopentyl GGAACACGTC
Dec. 4141 Pyropho. 4201	arboxylase TCTACCAAAG sphate Isom ATTTTATTGA	GAATAACTGC	AGCCCGGGAG	AAACGAATCT RBS GAGGATTACT GGTACGCTGG	TTGATTGACG Star ATATGCAAAC AAAAGTATGC	CAAAGACTGG rt Isopentyl GGAACACGTC CGCACACACG
Dec. 4141 Pyropho. 4201 4261	arboxylase TCTACCAAAG sphate Isom ATTTTATTGA GCAGACACCC	GAATAACTGC erase (idi) ATGCACAGGG	AGCCCGGGAG AGTTCCCACG CGCGTTCTCC	AAACGAATCT RBS GAGGATTACT GGTACGCTGG AGTTGGCTGT	TTGATTGACG Star ATATGCAAAC AAAAGTATGC TTAATGCCAA	CAAAGACTGG rt Isopentyl GGAACACGTC CGCACACACG AGGACAATTA
Dec. 4141 Pyropho. 4201 4261 4321	arboxylase TCTACCAAAG sphate Isom ATTTTATTGA GCAGACACCC TTAGTTACCC	GAATAACTGC erase (idi) ATGCACAGGG GCTTACATCT	AGCCCGGGAG AGTTCCCACG CGCGTTCTCC GAGCAAAAAA	AAACGAATCT RBS GAGGATTACT GGTACGCTGG AGTTGGCTGT GCATGGCCTG	TTGATTGACG Star ATATGCAAAC AAAAGTATGC TTAATGCCAA GCGTGTGGAC	CAAAGACTGG rt Isopentyl GGAACACGTC CGCACACACG AGGACAATTA TAACTCGGTT
Dec. 4141 Pyropho. 4201 4261 4321 4381	arboxylase TCTACCAAAG sphate Isom ATTTATTGA GCAGACACCC TTAGTTACCC TGTGGGCACC	GAATAACTGC erase (idi) ATGCACAGGG GCTTACATCT GCCGCGCACT	AGCCCGGGAG AGTTCCCACG CGCGTTCTCC GAGCAAAAAA AGAAAGCAAC	AAACGAATCT RBS GAGGATTACT GGTACGCTGG AGTTGGCTGT GCATGGCCTG GAAGACGCAG	TTGATTGACG Star ATATGCAAAC AAAAGTATGC TTAATGCCAA GCGTGTGGAC TGATCCGCCG	CAAAGACTGG rt Isopentyl GGAACACGTC CGCACACACG AGGACAATTA TAACTCGGTT TTGCCGTTAT
Dec. 4141 Pyropho. 4201 4261 4321 4381 4441	arboxylase TCTACCAAAG sphate Isom ATTTATTGA GCAGACACCC TTAGTTACCC TGTGGGCACC	GAATAACTGC erase (idi) ATGCACAGGG GCTTACATCT GCCGCGCACT CACAACTGGG	AGCCCGGGAG AGTTCCCACG CGCGTTCTCC GAGCAAAAAA AGAAAGCAAC GCCTCCTGAA	AAACGAATCT RBS GAGGATTACT GGTACGCTGG AGTTGGCTGT GCATGGCCTG GAAGACGCAG TCTATCTATC	TTGATTGACG Star ATATGCAAAC AAAAGTATGC TTAATGCCAA GCGTGTGGAC TGATCCGCCG CTGACTTTCG	CAAAGACTGG rt Isopentyl GGAACACGTC CGCACACACG AGGACAATTA TAACTCGGTT TTGCCGTTAT CTACCGCGCC
Dec. 4141 Pyropho. 4201 4261 4321 4381 4441 4501	arboxylase TCTACCAAAG sphate Isom ATTTATTGA GCAGACACCC TTAGTTACCC TGTGGGCACC GAGCTTGGCG ACCGATCCGA	GAATAACTGC erase (idi) ATGCACAGGG GCTTACATCT GCCGCGCACT CACAACTGGG TGGAAATTAC	AGCCCGGGAG AGTTCCCACG CGCGTTCTCC GAGCAAAAA AGAAAGCAAC GCCTCCTGAA GGAAAATGAA	AAACGAATCT RBS GAGGATTACT GGTACGCTGG AGTTGGCTGT GCATGGCCTG GAAGACGCAG TCTATCTATC GTGTGTCCGG	Star ATATGCAAAC AAAAGTATGC TTAATGCCAA GCGTGTGGAC TGATCCGCCG CTGACTTTCG TATTTGCCGC	CAAAGACTGG rt Isopentyl GGAACACGTC CGCACACACG AGGACAATTA TAACTCGGTT TTGCCGTTAT CTACCGCGCC ACGCACCACT
Dec. 4141 Pyropho. 4201 4261 4321 4381 4441 4501 4561	arboxylase TCTACCAAAG sphate Isom ATTTATTGA GCAGACACCC TTAGTTACCC TGTGGGCACC GAGCTTGGCG ACCGATCCGA AGTGCGTTAC	GAATAACTGC erase (idi) ATGCACAGGG GCTTACATCT GCCGCGCACT CACAACTGGG TGGAAATTAC GTGGCATTGT	AGCCCGGGAG AGTTCCCACG CGCGTTCTCC GAGCAAAAA AGAAAGCAAC GCCTCCTGAA GGAAAATGAA TGATGAAGTG	AAACGAATCT RBS GAGGATTACT GGTACGCTGG AGTTGGCTGT GCATGGCCTG GAAGACGCAG TCTATCTATC GTGTGTCCGG ATGGATTATC	Star ATATGCAAAC AAAAGTATGC TTAATGCCAA GCGTGTGGAC TGATCCGCCG CTGACTTTCG TATTTGCCGC AATGGTGTGA	CAAAGACTGG rt Isopentyl GGAACACGTC CGCACACACG AGGACAATTA TAACTCGGTT TTGCCGTTAT CTACCGCGCC ACGCACCACT TTTAGCAGAT
Dec. 4141 Pyropho. 4201 4261 4321 4381 4441 4501 4561	arboxylase TCTACCAAAG sphate Isom ATTTATTGA GCAGACACCC TTAGTTACCC TGTGGGCACC GAGCTTGGCG ACCGATCCGA AGTGCGTTAC	GAATAACTGC erase (idi) ATGCACAGGG GCTTACATCT GCCGCGCACT CACAACTGGG TGGAAATTAC GTGGCATTGT AGATCAATGA	AGCCCGGGAG AGTTCCCACG CGCGTTCTCC GAGCAAAAA AGAAAGCAAC GCCTCCTGAA GGAAAATGAA TGATGAAGTG	AAACGAATCT RBS GAGGATTACT GGTACGCTGG AGTTGGCTGT GCATGGCCTG GAAGACGCAG TCTATCTATC GTGTGTCCGG ATGGATTATC	Star ATATGCAAAC AAAAGTATGC TTAATGCCAA GCGTGTGGAC TGATCCGCCG CTGACTTTCG TATTTGCCGC AATGGTGTGA	CAAAGACTGG rt Isopentyl GGAACACGTC CGCACACACG AGGACAATTA TAACTCGGTT TTGCCGTTAT CTACCGCGCC ACGCACCACT TTTAGCAGAT
Dec. 4141 Pyropho. 4201 4261 4321 4381 4441 4501 4561	arboxylase TCTACCAAAG sphate Isom ATTTATTGA GCAGACACCC TTAGTTACCC TGTGGGCACC GAGCTTGGCG ACCGATCCGA AGTGCGTTAC	GAATAACTGC erase (idi) ATGCACAGGG GCTTACATCT GCCGCGCACT CACAACTGGG TGGAAATTAC GTGGCATTGT AGATCAATGA	AGCCCGGGAG AGTTCCCACG CGCGTTCTCC GAGCAAAAA AGAAAGCAAC GCCTCCTGAA GGAAAATGAA TGATGAAGTG CACGCCGTGG	AAACGAATCT RBS GAGGATTACT GGTACGCTGG AGTTGGCTGT GCATGGCCTG GAAGACGCAG TCTATCTATC GTGTGTCCGG ATGGATTATC GCGTTCAGTC	TTGATTGACG Star ATATGCAAAC AAAAGTATGC TTAATGCCAA GCGTGTGGAC TGATCCGCCG CTGACTTTCG TATTTGCCGC AATGGTGTGA CGTGGATGGT	CAAAGACTGG rt Isopentyl GGAACACGTC CGCACACACG AGGACAATTA TAACTCGGTT TTGCCGTTAT CTACCGCGCC ACGCACCACT TTTAGCAGAT GATGCAGGCG

	RBS Start Farnesyl
	4741 TCCACTAGTT CTAGAGCGGC CGCCACCGCG GAGGAGGAAT GAGTAATGGA CTTTCCGCAG
	Pyrophosphate Synthase (ispA)
5	4801 CAACTCGAAG CCTGCGTTAA GCAGGCCAAC CAGGCGCTGA GCCGTTTTAT CGCCCCACTG
	4861 CCCTTTCAGA ACACTCCCGT GGTCGAAACC ATGCAGTATG GCGCATTATT AGGTGGTAAG
	4921 CGCCTGCGAC CTTTCCTGGT TTATGCCACC GGTCATATGT TCGGCGTTAG CACAAACACG
	4981 CTGGACGCAC CCGCTGCCGC CGTTGAGTGT ATCCACGCTT ACTCATTAAT TCATGATGAT
	5041 TTACCGGCAA TGGATGATGA CGATCTGCGT CGCGGTTTGC CAACCTGCCA TGTGAAGTTT
10	5101 GGCGAAGCAA ACGCGATTCT CGCTGGCGAC GCTTTACAAA CGCTGGCGTT CTCGATTTTA
	5161 AGCGATGCCG ATATGCCGGA AGTGTCGGAC CGCGACAGAA TTTCGATGAT TTCTGAACTG
	5221 GCGAGCGCCA GTGGTATTGC CGGAATGTGC GGTGGTCAGG CATTAGATTT AGACGCGGAA
	5281 GGCAAACACG TACCTCTGGA CGCGCTTGAG CGTATTCATC GTCATAAAAC CGGCGCATTG
15	5341 ATTCGCGCCG CCGTTCGCCT TGGTGCATTA AGCGCCGGAG ATAAAGGACG TCGTGCTCTG
15 宣	5401 CCGGTACTCG ACAAGTATGC AGAGAGCATC GGCCTTGCCT TCCAGGTTCA GGATGACATC
TOTAL	5461 CTGGATGTGG TGGGAGATAC TGCAACGTTG GGAAAACGCC AGGGTGCCGA CCAGCAACTT
I STATE OF THE STA	5521 GGTAAAAGTA CCTACCCTGC ACTTCTGGGT CTTGAGCAAG CCCGGAAGAA AGCCCGGGAT
To the second se	5581 CTGATCGACG ATGCCCGTCA GTCGCTGAAA CAACTGGCTG AACAGTCACT CGATACCTCG
≅ ≅ =	
20	Stop Farnesyl Pyrophosphate Synthase
and the state of t	5641 GCACTGGAAG CGCTAGCGGA CTACATCATC CAGCGTAATA AATAAGAGCT CCAATTCGCC
plants que	5701 CTATAGTGAG TCGTATTACG CGCGCTCACT GGCCGTCGTT TTACAACGTC GTGACTGGGA
	5761 AAACCCTGGC GTTACCCAAC TTAATCGCCT TGCAGCACAT CCCCCTTTCG CCAGCTGGCG
•	5821 TAATAGCGAA GAGGCCCGCA CCGATCGCCC TTCCCAACAG TTGCGCAGCC TGAATGGCGA
25	5881 ATGGAAATTG TAAGCGTTAA TATTTTGTTA AAATTCGCGT TAAATTTTTG TTAAATCAGC

5941 TCATTTTTTA ACCAATAGGC CGA